

## East meets West or Out of Africa? Genomic evidence and trade data to understand pest risks and globalization of *Spodoptera frugiperda*

WT Tay, D Kunz, T Walsh, S. Elfekih, S Downes, A Padovan, D Kriticos, K Nam, E d'Alençon, N Nègre, Y Wu, J Zhang, C Czepak, M Otim, K Gordon

33-6 Sept 2019 | iPRRG 2019 Annual Meeting | Poznań, Poland

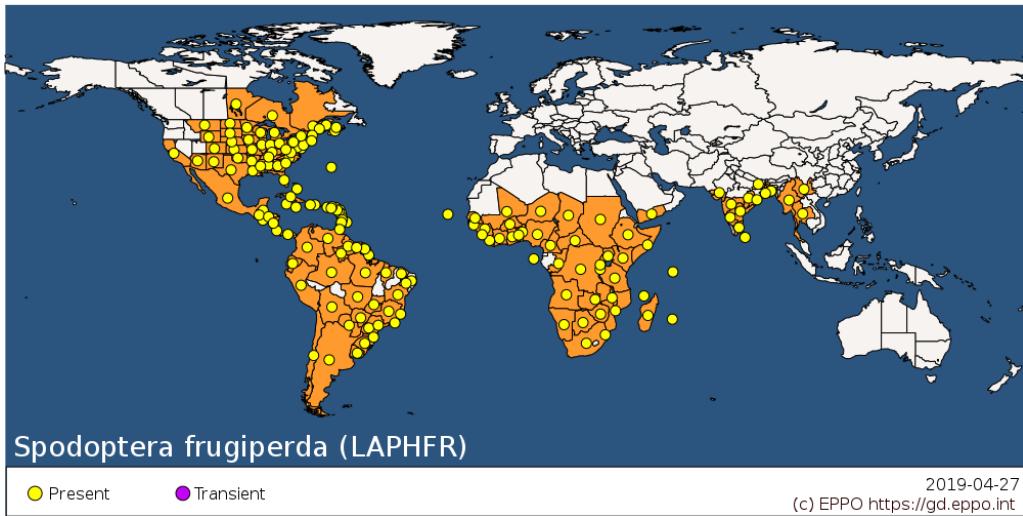
CSIRO HEALTH & BIOSECURITY | RISK EVALUATION & PREPAREDNESS PROGRAM  
[www.csiro.au](http://www.csiro.au)



**INRA**  
SCIENCE & IMPACT



# *Spodoptera frugiperda*: Africa & beyond



Food and Agriculture Organization of the United Nations

BRIEFING NOTE ON FAO ACTIONS  
ON FALL ARMYWORM

FEED THE FUTURE  
The U.S. Government's Global Hunger & Food Security Initiative



Figure 1 Distribution map of *Spodoptera frugiperda* from the EPPO Global Database accessed 27/04/2019.

**TABLE 1: TOP 20 PESTS BY RECENT PUBLICATION NUMBERS**

Ranking is based on abstract searches on all 1,187 arthropod pests with full datasheets in the CABI Compendia on crop protection, forestry and invasive species. Also included are the number of pesticide active ingredients to which they are reported to be resistant, and summaries of their reported distribution and reported hosts.

| SCIENTIFIC NAME                 | COMMON NAME      | NUMBER OF PUBLICATIONS (2012-16) | NUMBER OF PESTICIDES RESISTANT TO | CONTINENTS PRESENT <sup>a</sup>               | NUMBER OF HOST GENERA <sup>b</sup> | TOP HOSTS (BY PUBLICATION NUMBER) <sup>c*</sup>                |
|---------------------------------|------------------|----------------------------------|-----------------------------------|---|------------------------------------|--|
| 5 <i>Spodoptera litura</i>      | taro caterpillar | 853                              | 39                                | Asia, Africa, N America, Europe, Oceania *    | 66                                 | <i>Glycine max</i> (soybean), <i>Arachis hypogaea</i> (peanut) |
| 8 <i>Spodoptera frugiperda</i>  | fall armyworm    | 668                              | 24                                | Africa, N/S America, Europe (few occurrences) | 72                                 | <i>Zea mays</i> (maize), <i>Gossypium</i> (cotton)             |
| 11 <i>Spodoptera exigua</i>     | beet armyworm    | 506                              | 39                                | Asia, Africa, N America, Europe, Oceania      | 50                                 | <i>Gossypium</i> (cotton), <i>Zea mays</i> (maize)             |
| 16 <i>Spodoptera littoralis</i> | cotton leafworm  | 401                              | 30                                | Asia, Africa, Europe                          | 96                                 | <i>Gossypium</i> (cotton), <i>Zea mays</i> (maize)             |



*Spodoptera frugiperda*

(Fall Armyworm)

Hannalene du Plessis<sup>1</sup>, Johnnie van den Berg<sup>1</sup>, Noboru Ota<sup>2</sup> & Darren J. Kriticos<sup>2,3,\*</sup><sup>1</sup>Unit for Environmental Sciences and Management, North-West University, Potchefstroom, South Africa<sup>2</sup>CSIRO, GPO Box 1700, Canberra, Australia<sup>3</sup>The University of Queensland

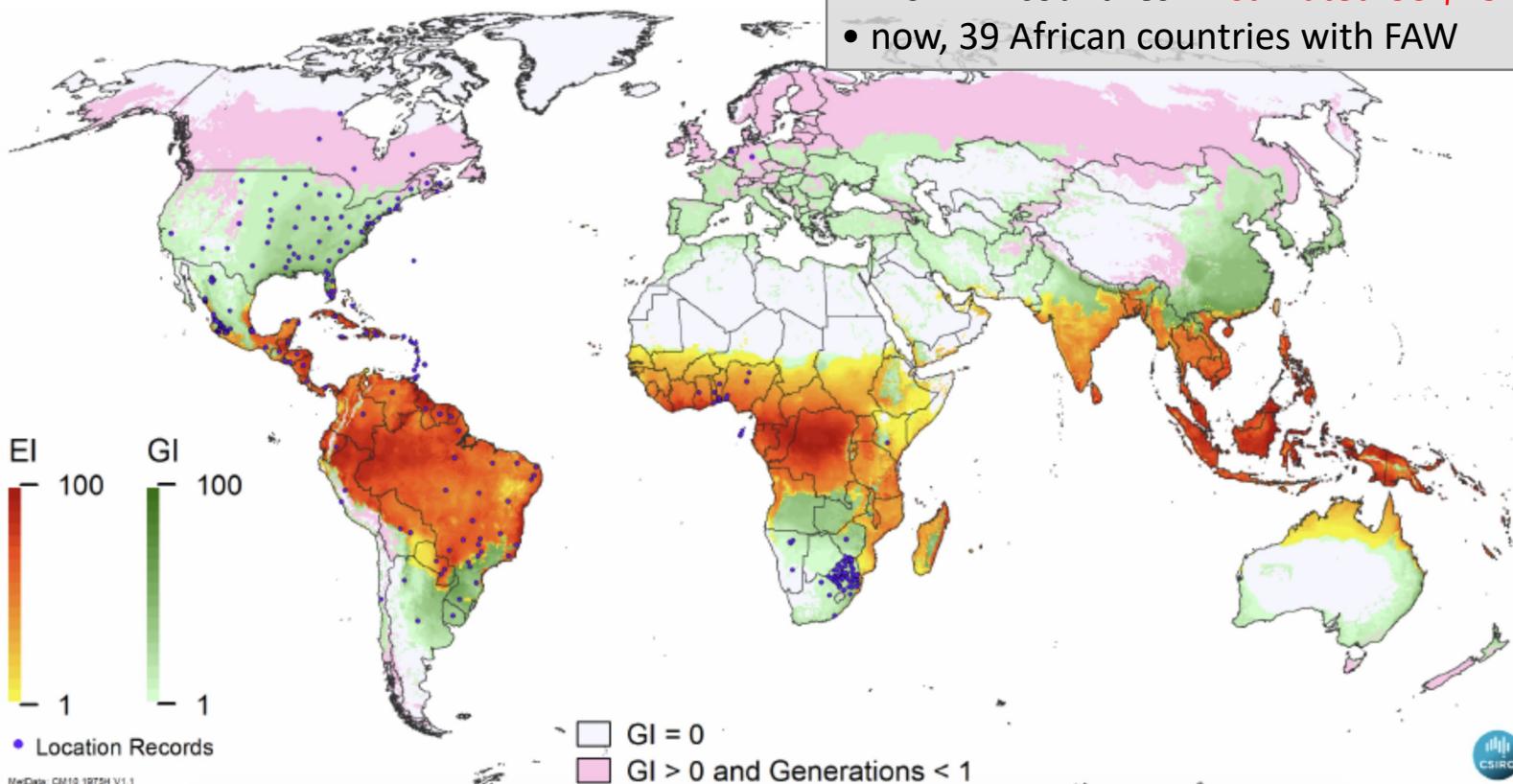
\*Correspondence: Darren.Kriticos@csiro.au

## Fall Armyworm Status

Impacts and control options in Africa:  
Preliminary Evidence Note (April 2017)P. Abrahams, T. Beale, M. Cock, N. Corniani, R. Day\*, J. Godwin,  
S. Murphy, G. Richards & J. Vos

| FAW affected crops<br>in all countries | Total production (tonnes, m)<br>assuming no FAW | Yield loss<br>(tonnes, m) | Estimated/predicted loss<br>(US\$, m) |
|--|---|---------------------------|---------------------------------------|
| Maize                                  | 67.0  | 13.5                      | 3,058                                 |
| Sorghum                                | 25.5  | 1.90                      | 827                                   |
| Rice, paddy                            | 17.1  | 9.6                       | 6,699                                 |
| Sugarcane                              | 90.1  | 46.0                      | 2,798                                 |
| <b>Total</b>                           |   |                           | <b>13,383</b>                         |

- From 11 countries... Estimated US \$13.3 billion
- now, 39 African countries with FAW



# The Very Hungry Caterpillar ...

 ANTARANEWS.COM

**Ministry takes precautionary measures against armyworm pest attacks**

© 31st May 2019



 KHMER TIMES

National June 15, 2019

**Worm epidemic destroys 10, 000 hectares of corn**

Mom Kunthea / Khmer Times  
Share: f t G+ D



The Fall Armyworm which ravaged corn crops in South

 nature International journal of science

NEWS • 17 JUNE 2019

Caterpillar's devastating march across China spurs hunt for native predator

Scientists hope that insects such as stink bugs could keep the fall armyworm in check.

Fall armyworms have come to Nepal from America, and they can trouble maize farmers here

 Bhutan



Adults reared from larvae  
(T. Zangpo, Dept. Ag.)

 XINHUANET

Crop-eating fall armyworm spread across Taiwan

Source: Xinhua | 2019-06-18 16:56:45 |

f t in s w

TAIPEI, June 18 (Xinhua) -- The fall armyworm, a crop-eating pest, has been detected in 18 counties and cities across Taiwan as of Monday, the island's agricultural authorities said.

 malaysiakini

Home News Parliament Editor's Pick Sp



Fall armyworm invades crops across Asia

Patpicha Tanakasempipat and Naveen Thukral, Reuters  
Jun 20th, 2019 (Updated Jun 20th, 2019)

- Confusing species status: corn- or rice-preferred ...
- Highly invasive: detected in > 60 countries since 2016
- African + SE Asia: corn yield loss: 50%
- China: 2<sup>nd</sup> world corn producer
- How did it spread across the world?
  - Origins of invasive Sfc & Sfr populations?
  - Corn / Rice; hybrids?

# Global effort on forecasting FAW spread

NeoBiota 40: 25–50 (2018)  
doi: 10.3897/neobiota.40.28165  
<http://neobiota.pensoft.net>

RESEARCH ARTICLE



## Forecasting the global extent of invasion of the cereal pest *Spodoptera frugiperda*, the fall armyworm

Regan Early<sup>1</sup>, Pablo González-Moreno<sup>2</sup>, Sean T. Murphy<sup>2</sup>, Roger Day<sup>3</sup>

<sup>1</sup> Centre for Ecology and Conservation, Exeter University, Penryn, Cornwall, TR10 9FE, UK <sup>2</sup> CABI, Bawdsey Lane, Egham TW20 9TY, UK <sup>3</sup> CABI, Canary Bird, 673 Limuru Road, Muthaiga, PO Box 633-00621, Nairobi, Kenya

- trade & transport routes from Africa are important factors

- high threat of FAW originating from Africa to:  
**China, Thailand, Australia, Indonesia, Malaysia, Philippines**

### • Genomic evidence?



Food and Agriculture Organization  
of the United Nations

Fall Armyworm likely to spread from India to other parts of Asia with South East Asia and South China most at risk

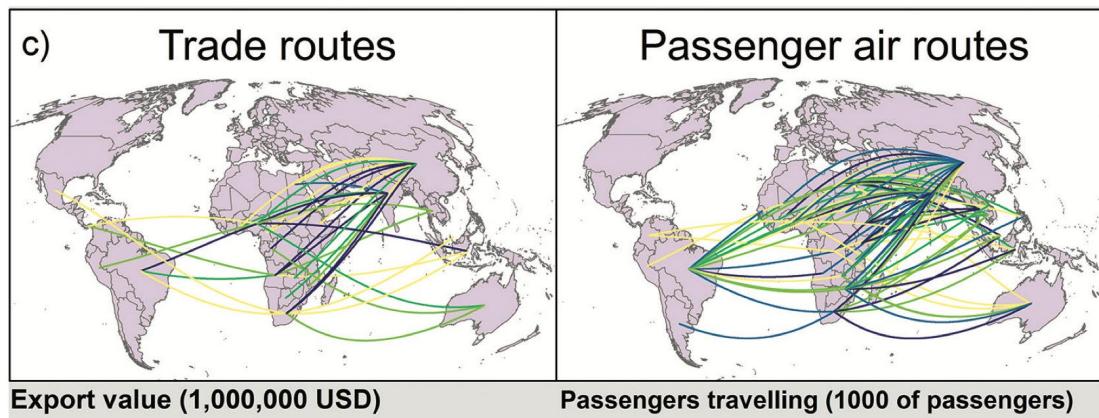
FAO offers its expertise to farmers and governments in Asia on best ways to manage the invasive pest

© FAO, 2019

- Spread of FAW **likely** from India to South China and South East Asia

### • Genomic evidence?

- UN data (total exports) from sub-Saharan African countries to all countries between 2012-2016.
- total trade volume >USD\$ 500 M as most likely to introduce FAW
- Passenger (air) also considered



## Molecular identification of invasive fall armyworm *Spodoptera frugiperda* in Yunnan Province

ZHANG Lei<sup>1</sup>, JIN Minghui<sup>1</sup>, ZHANG Dandan<sup>2</sup>, JIANG Yuying<sup>2</sup>, LIU Jie<sup>3</sup>, WU Kongming<sup>2</sup>, XIAO Yutao<sup>1</sup>  
 1. Agricultural Genomics Institute at Shenzhen, Chinese Academy of Agricultural Sciences, Shenzhen 518120, China;  
 2. State Key Laboratory for Biology of Plant Diseases and Insect Pests, Institute of Plant Protection, Chinese Academy of Agricultural Sciences, Beijing 100193, China; 3. National Agro-Tech Extension and Service Center, Beijing 100125, China

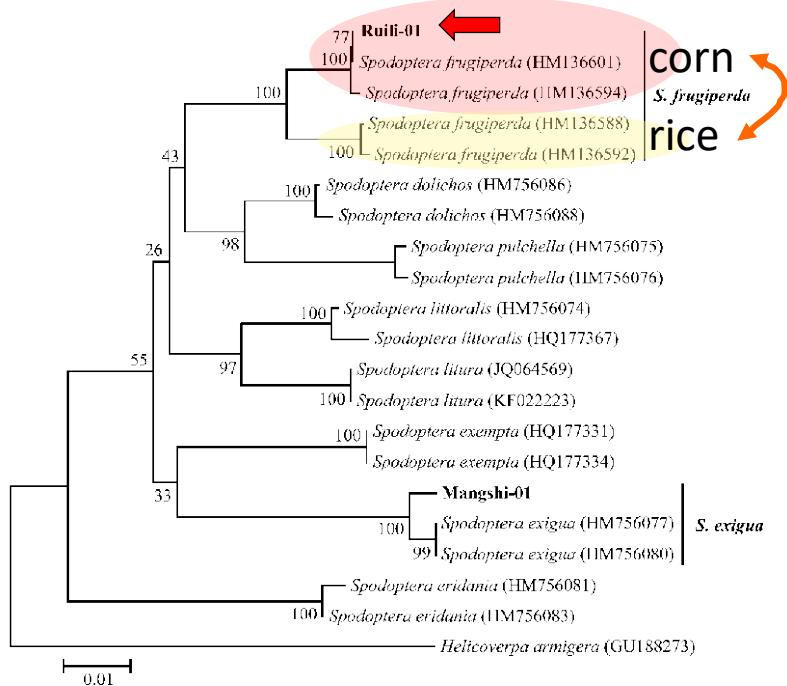


图 2 基于 CO I 序列构建的 NJ 系统树

Fig. 2 Phylogenetic tree constructed by NJ analysis based on CO I gene fragments

K2P, NJ, 1,000 bootstrap replications

Genetic identity of Myanmar populations?

## Analysis of migration routes of the fall armyworm *Spodoptera frugiperda* (J. E. Smith) from Myanmar to China

WU Qitun<sup>1</sup>, JIANG Yuying<sup>2</sup>, WU Kongming<sup>1</sup>

1. State Key Laboratory for Biology of Plant Diseases and Insect Pests, Institute of Plant Protection, Chinese Academy of Agricultural Sciences, Beijing 100193, China; 2. National Agro-Tech Extension and Service Center, Beijing 100125, China

By end of 2018, Myanmar FAW “sporadically entered southwestern Yunnan”

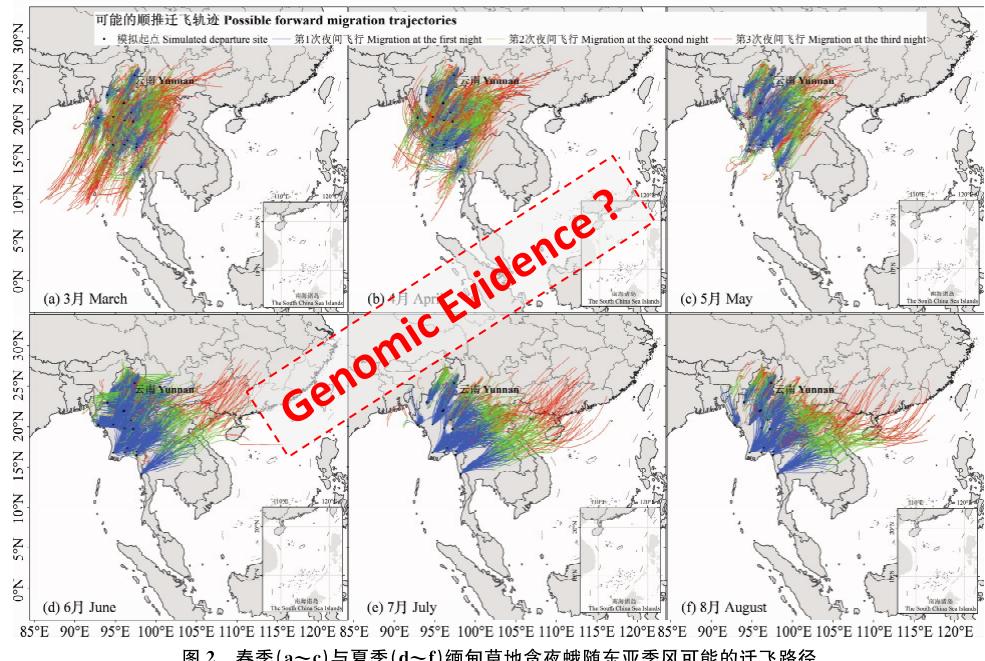


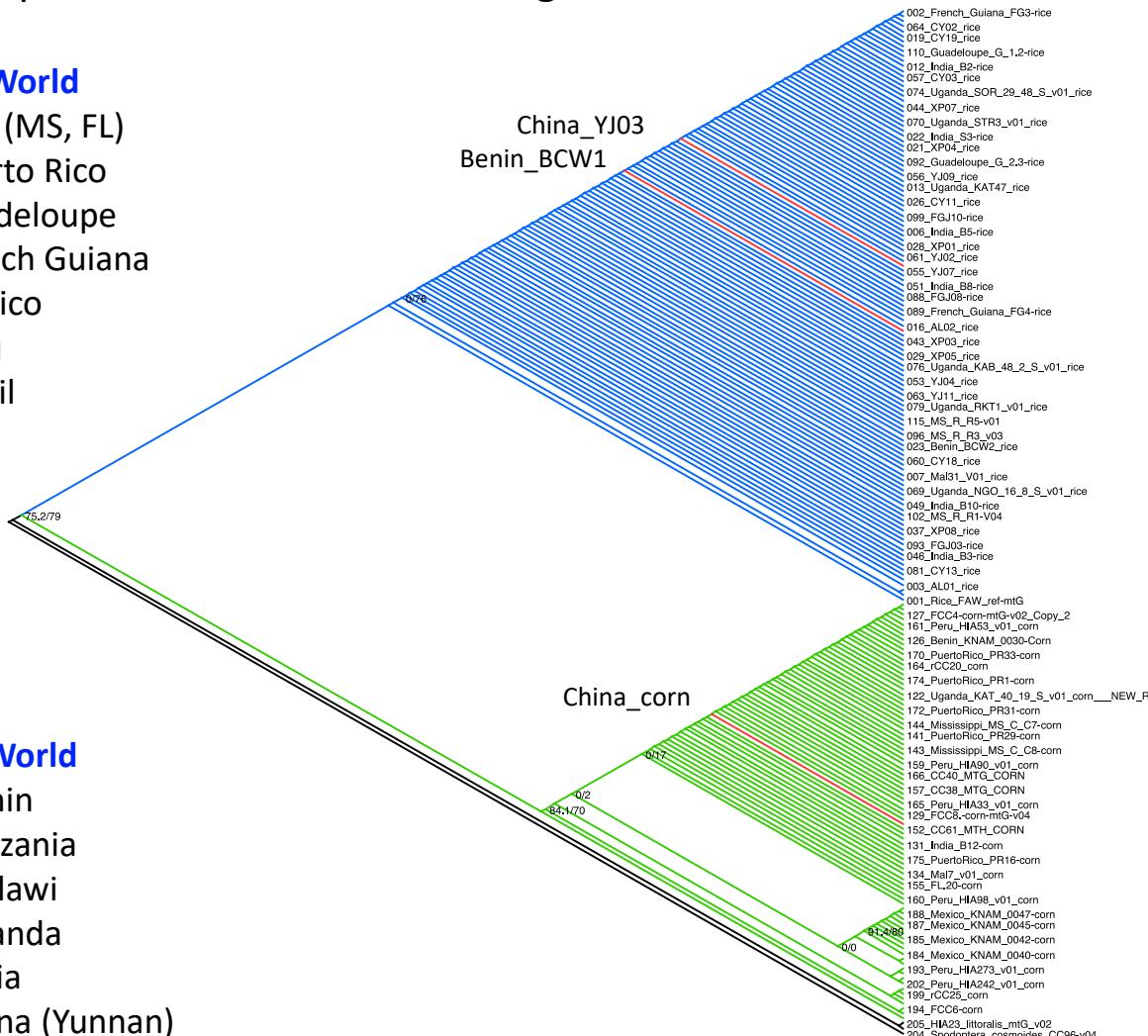
图 2 春季(a~c)与夏季(d~f)缅甸草地贪夜蛾随东亚季风可能的迁飞路径

Fig. 2 Possible migration trajectories of the fall armyworm in Myanmar with East Asian monsoon in spring (a~c) and summer (d~f)

# 638bp standard mtCOI barcode gene

## New World

- USA (MS, FL)
- Puerto Rico
- Guadeloupe
- French Guiana
- Mexico
- Peru
- Brazil



• Single origins for Sfc & Sfr...

Sfr

100% nucleotide identity  
within the major clade

All invasive + New World  
populations

Sfc

100% nucleotide identity  
within the major clade

All invasive + New World  
(excludes MEX) populations

Despite extensive sampling of native populations, failed to ascertain source populations!

# Population origins?

Global populations whole genome sequenced

- USA (MS, FL)
- Puerto Rico, Guadeloupe
- Brazil, Peru, Mexico, French Guiana
- E. Africa (Tz, Ug, Mw)
- W. Africa (Benin)
- India
- China

Sfr: N = 120

Sfc: N = 84

*S. eridania*

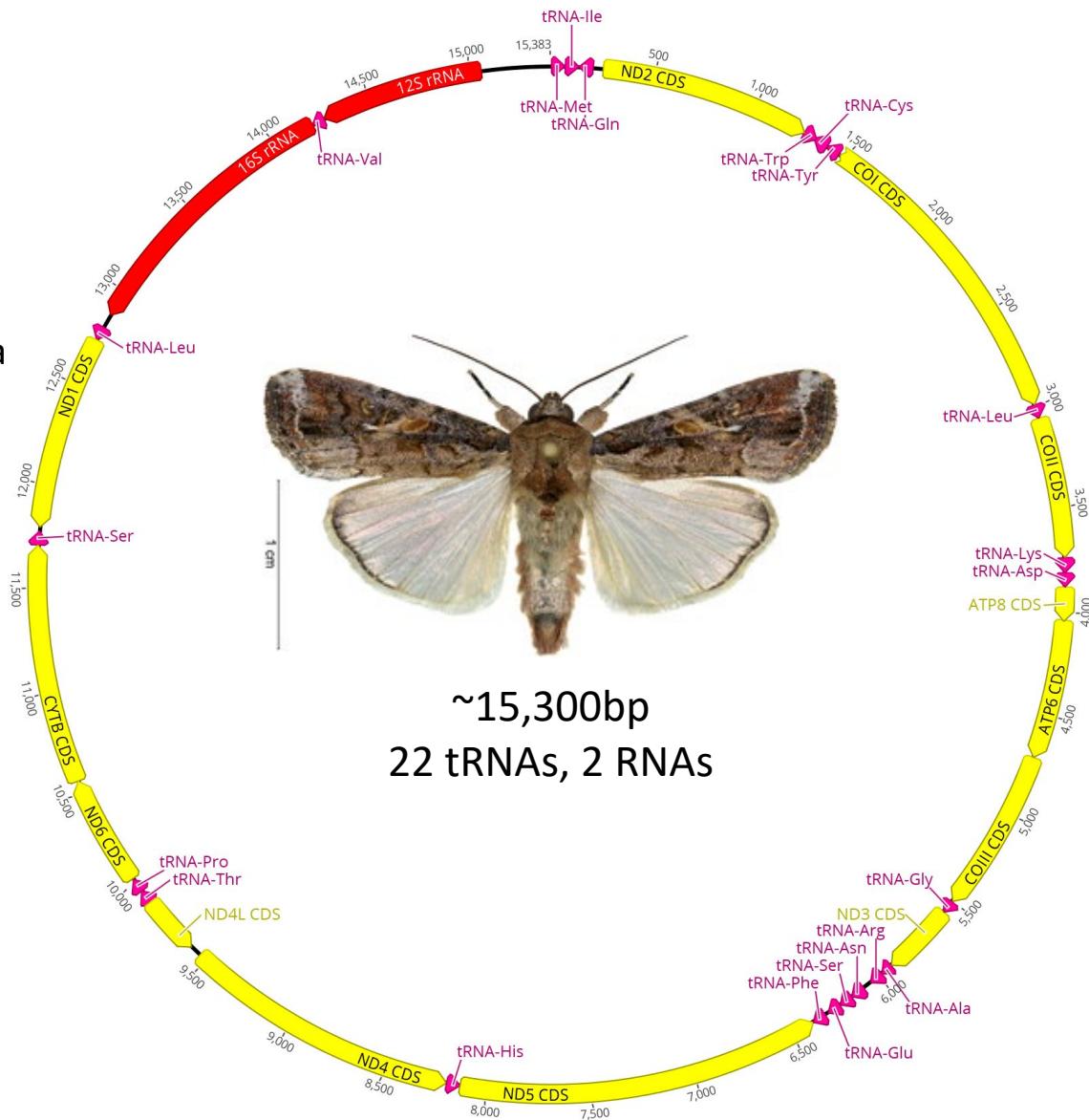
*S. litura*

*S. littoralis*

} ~ 3.16 Mbp

## Missing:

- SE Asia, Japan, South Korea
- West Africa
- South Africa
- Middle East



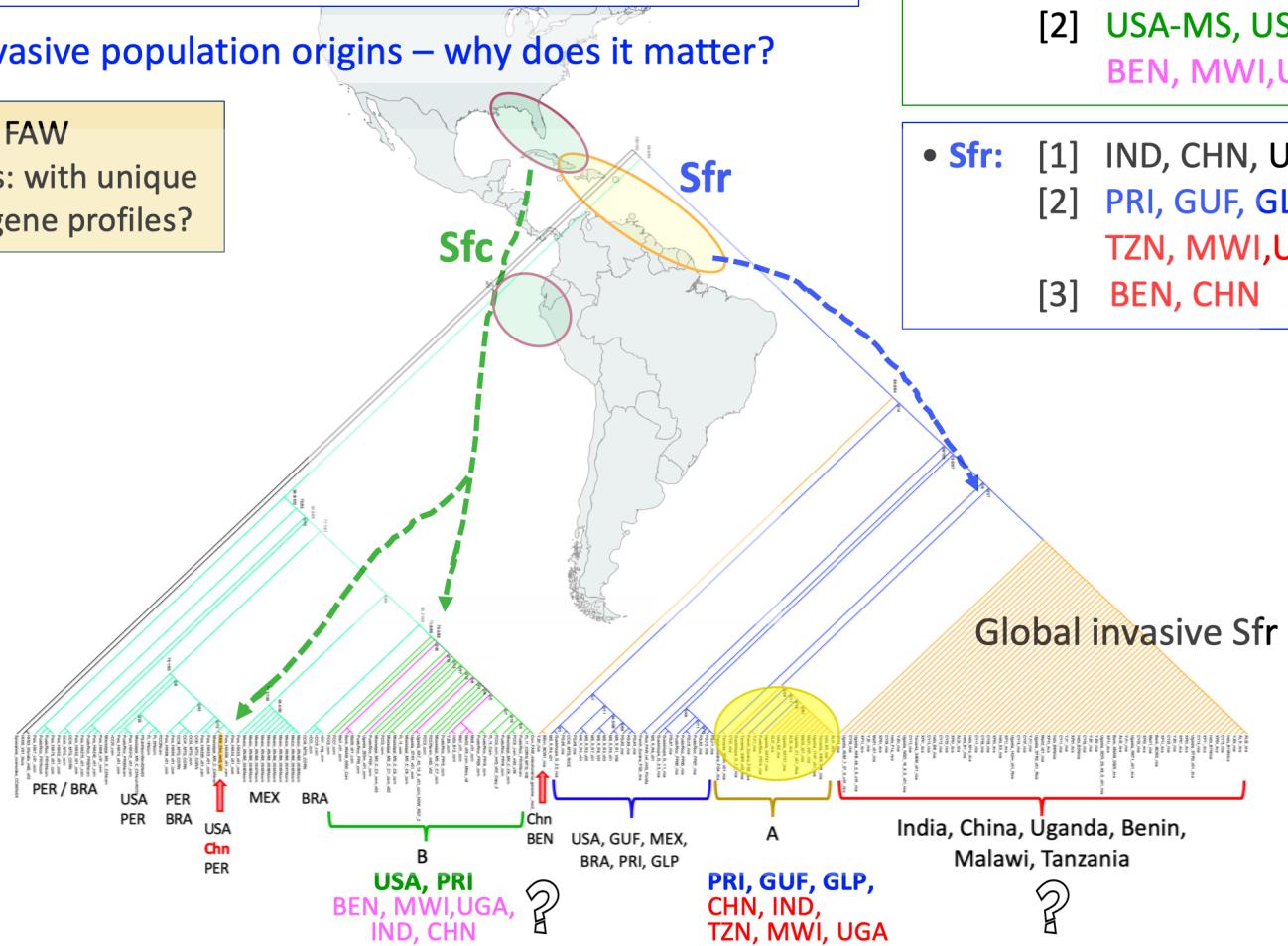


## Going global – genomic insights into insect invasions

Wee Tek Tay and Karl Heinrich Julius Gordon

### Tracing Invasive population origins – why does it matter?

New World FAW populations: with unique resistance gene profiles?



- Evidence of multiple introductions:

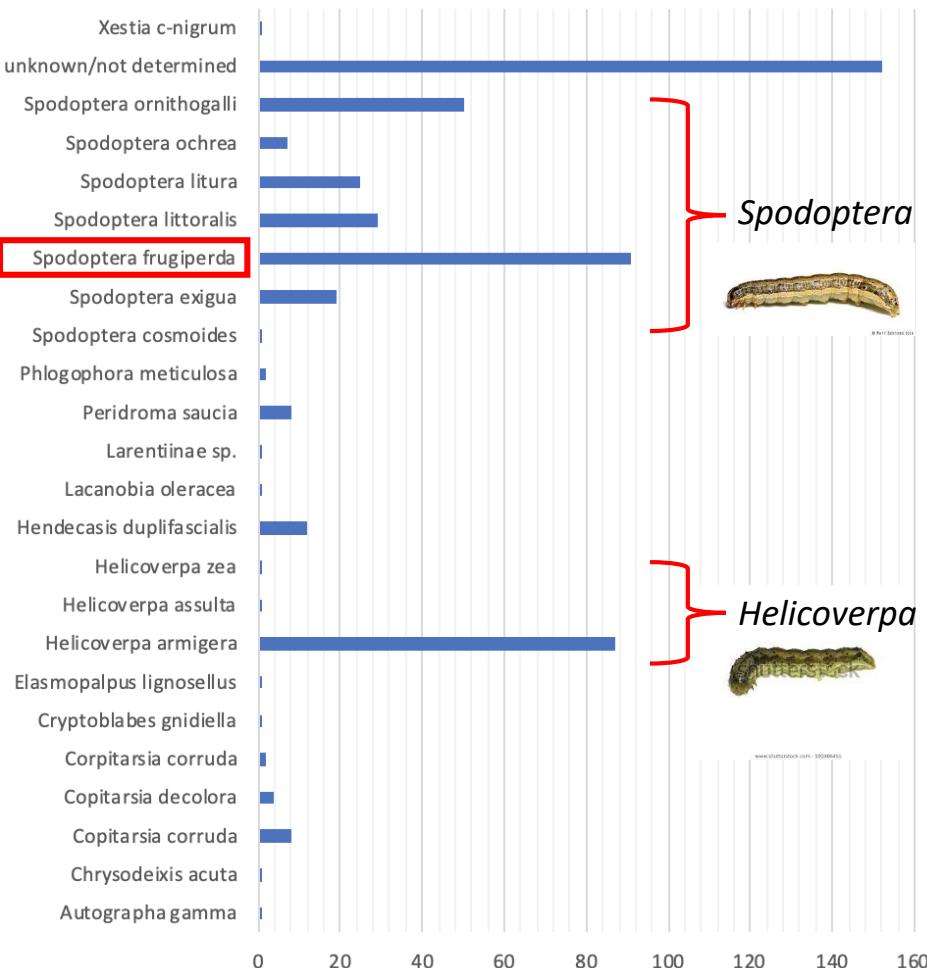
- Sfc:**
  - [1] USA-MS, CHN, PER
  - [2] USA-MS, USA-FL, PRI, BEN, MWI, UGA, IND, CHN

- Sfr:**
  - [1] IND, CHN, UGA, BEN, MWI, TZN
  - [2] PRI, GUF, GLP, TZN, MWI, UGA, IND, CHN
  - [3] BEN, CHN

## Australia pre-border interceptions (2016 – 2019)

| Region        | <i>Helicoverpa</i> | <i>Spodoptera</i> | others     |
|---------------|--------------------|-------------------|------------|
| Asia          | 38                 | 46                | 48         |
| Oceania       | 0                  | 3                 | 2          |
| Africa        | 40                 | 16                | 45         |
| North America | 0                  | 1                 | 0          |
| South America | 9                  | 155               | 93         |
| Europe        | 2                  | 1                 | 7          |
| <b>total</b>  | <b>89</b>          | <b>222 *</b>      | <b>195</b> |

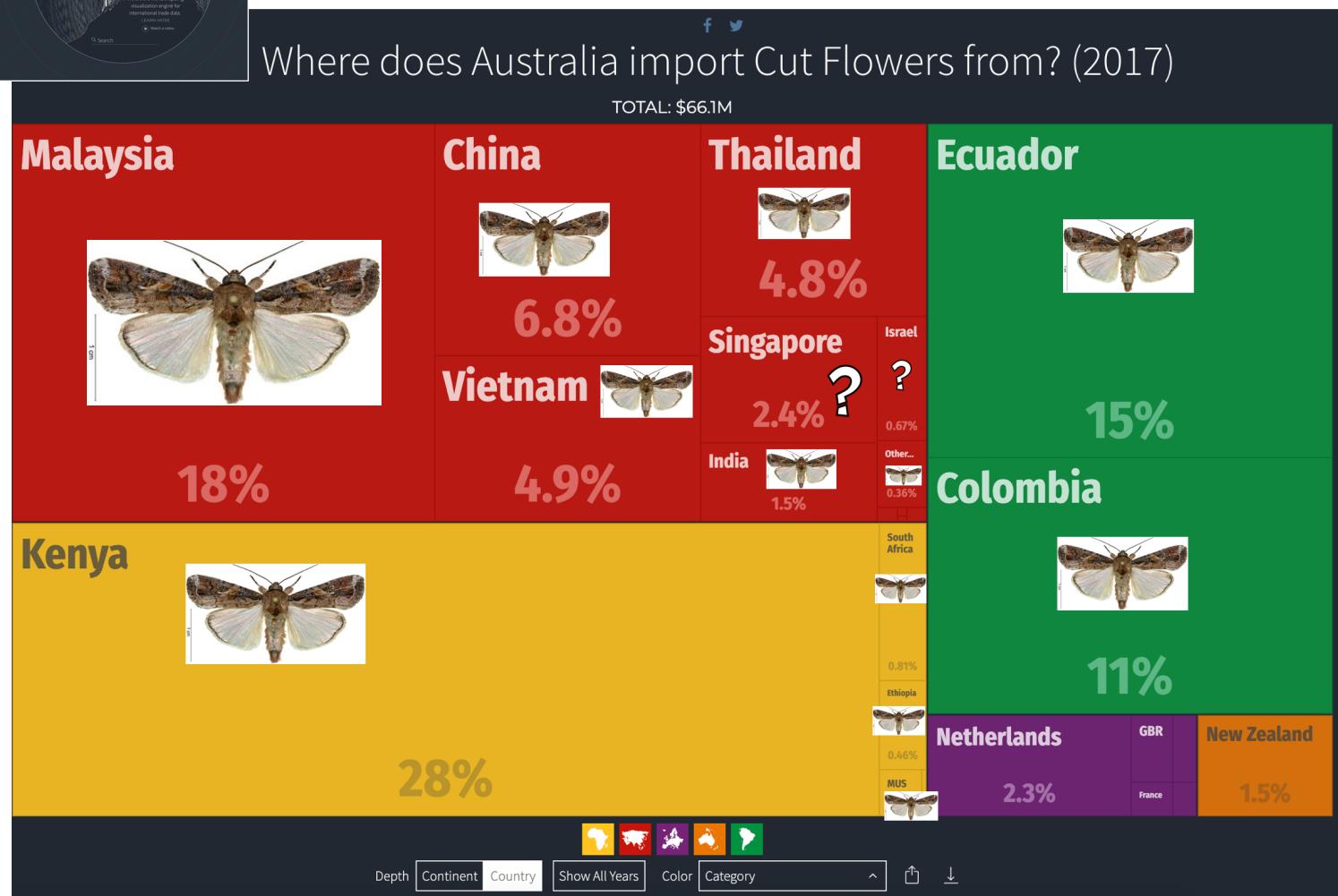
## total detection of invasive lepidopteran species



- FAW + other *Spodoptera* spp. intercepted
- FAW intercepted (Africa, Asia, Sth America)
- Cut flowers, fresh vegetables



# Risks to Australia's agriculture sector?

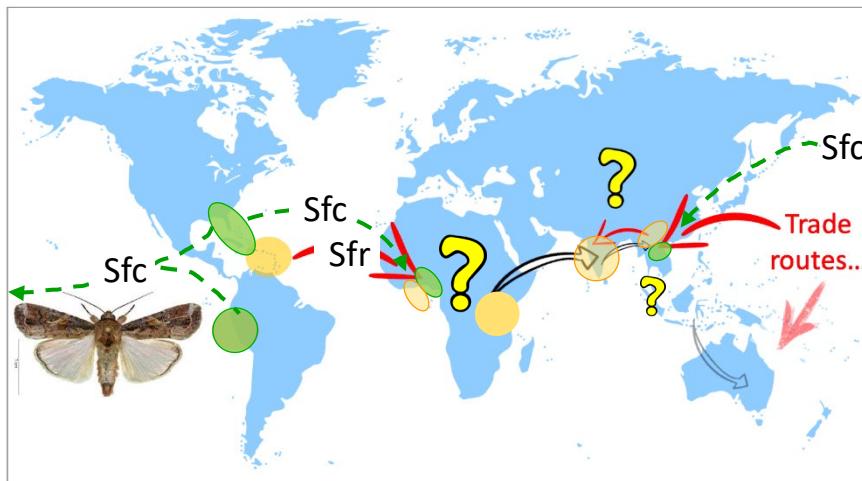


- Singapore
- Israel
- Malaysia
- South Korea
- Japan
- Hong Kong
- Nepal
- India
- Thailand
- Vietnam
- China
- Kenya
- South Africa
- Mauritius
- Ethiopia
- Ecuador
- Colombia

91.85%

# May be... Out of Africa & East meets West?

- High genetic diversity in African FAW:  $\geq 8$  maternal lineages
- Multiple introduction pathways into the Old World: human-assisted / international trade
- Spread across the Old World: Human-assisted / natural migration?



?

Implications on pest management strategies, esp. in S.E.A.

- *S. eridania*
- *S. orchae*
- *S. cosmoides*
- *S. ornithogalli*
- *S. marima*
- etc!!

- Multiple New World origins for Sfc and Sfr
- ‘African origin’ theory + ‘Myanmar source pop’ to China ...  
Unresolved by mitogenomes; genome-wide SNPs evidence needed!

**Key Message:** Global population genomics and trade pathway analyses needed to understand gene flow patterns and predict future pest spread.

# Acknowledgements



- CSIRO (H&B R-08681-1)
- International collaborators (UFG, NaCRRI), CSIRO colleagues



- INRA colleagues: Emmanuelle d'Alençon, Kiwoong Nam, Nicolas Nègre
- FAW samples: UFG, NaCRRI, DoA (Australia), INRA, TPRI (Dr Maneno Chidege, Tanzania), DARS (Dr Donald Kachigamba, Malawi)

Wee Tek Tay, PhD

t +61 2 6246 4286

e weetek.tay@csiro.au

w www.csiro.au/en/Research/BF