

How to analyze life table data with different number of stages

Prof. Dr. Hsin Chi

齊心教授

Department of Plant Production and Technologies

Faculty of Agricultural Sciences and Technologies

Ömer Halisdemir Üniversitesi, Niğde, Turkey

Visiting professor, Fujian Agriculture and Forestry University, China

Creative thinking

If individuals have different number of stages, e.g., some have 5 larval stages while some have 6 larval stages, but you want to show curves with L6, how can you analyze your data to show the survival curves? Or. How can you use the TWOSEX program to show the difference in stage numbers among individuals?

Can you do this? NO!

"Project: Scale insect at 25°C"

"User: X. X. Yang"

"Date: 2001.4.16.-2001.6.10."

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3,9

F,Egg,L1,L2,L3,L4,L5,L6,Pupa,Female

M,Egg,L1,L2,L3,L4,L5,L6,Pupa,Male

N,Egg,L1,L2,L3,L4,L5,L6,Unknown

L1,L6

1,M,7,2,5,6,4,**5,3**,12,30

2,M,7,2,4,4,4,**6,0**,10,44

3,N,7,-7

File A: Combine L5 and L6 to get population parameters and curves from Egg to L4

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3,**8**

F,Egg,L1,L2,L3,L4,**L5-6**,Pupa,Female

M,Egg,L1,L2,L3,L4,**L5-6**,Pupa,Male

N,Egg,L1,L2,L3,L4,**L5-6**,Pupa,Unknown

L1,**L5-6**

1,M,7,2,5,6,4,**8**,12,30

2,M,7,2,4,4,4,**6**,10,44

3,N,7,-7

File B: To get correct L1 to adult curves for L5 type insects: change all insects to N,-1 except those go through L5 and adult stage

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3,**8**

F,Egg,L1,L2,L3,L4,L5,Pupa,Female

M,Egg,L1,L2,L3,L4,L5,Pupa,Male

N,Egg,L1,L2,L3,L4,L5,Pupa,Unknown

L1,L5

1,N,-1

2,M,7,2,4,4,4,**6**,10,44

3,N,-1

File C: To get correct L1 to adult curves for L6 type insects: change all insects to N,-1 except those go through L6 and adult stage

"Project: Scale insect at 25C"

"User: X. X. Yang"

"Date: 2001.4.16.-2001.6.10."

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3,9

F,Egg,L1,L2,L3,L4,L5,L6,Pupa,Female

M,Egg,L1,L2,L3,L4,L5,L6,Pupa,Male

N,Egg,L1,L2,L3,L4,L5,L6,Pupa,Unknown

L1,L6

1,M,7,2,5,6,4,**5,3**,12,30

2,N,-1

3,N,-1

File D (or N5): To get correct L1 to pupa curves for N type insects died before L5, in L5 or Pupa (through L5): change all F, and M type, and those died in L6 and Pupa (through L6) to N,-1 and keep rest N type as they are.

"Project: Scale insect at 25C"

"User: X. X. Yang"

"Date: 2001.4.16.-2001.6.10." **55,N,-1**

50 **56,N,-1**

3,**8**

.....

L1,L5

1,N,-1

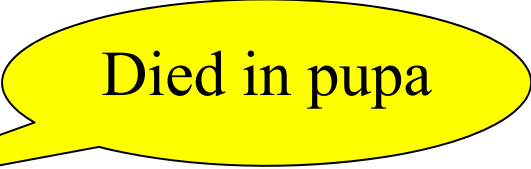
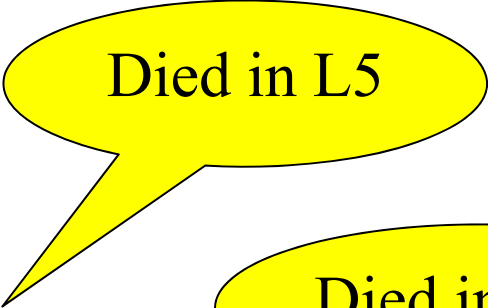
2,N,-1

3,N,7,-7

....

35,N,3,4,3,5,4,-5

36,N,3,3,5,6,4,5,-7



File E (or N6): To get correct L1 to pupa curves for N type insects died in L6 or Pupa (through L6): change all F, and M type, and those died in L5 and Pupa (through L5) to N,-1 and keep rest N type as they are.

"Project: Scale insect at 25C"

"User: X. X. Yang"

"Date: 2001.4.16.-2001.6.10." **55,N,3,4,4,5,4,5,-6**

50 **56,N,3,3,4,5,5,4,6,-5**

3,**9**

.....

L1,L6

1,N,-1

2,N,-1

3,N,-1

....

35,N,-1

36,N,-1

Died in L6

Died in pupa

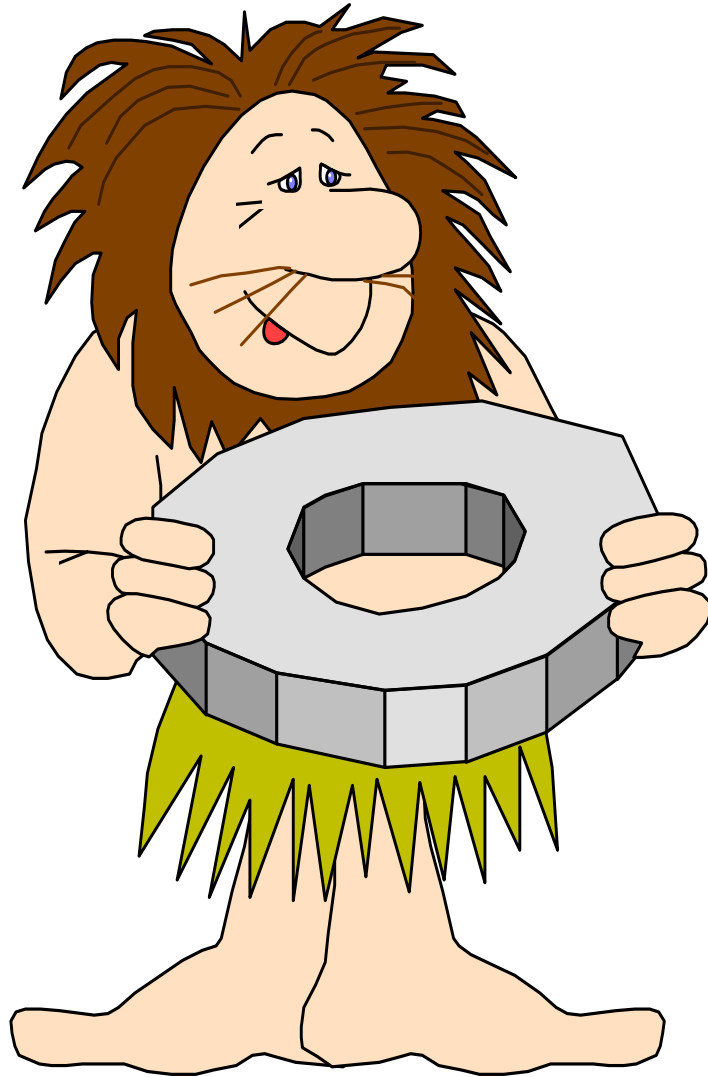
How to prepare figures

- Use file A to prepare all curves of s_{xj} (with L5+L6 combined).
- Import egg s_{xj} from file A, s_{xj} of L1-adult data of file B, s_{xj} of L1-adult data of file C, s_{xj} of L1-pupa of file D to a worksheet. Add all L1s to a new column (i.e., pooled-L1 and it should be exactly the same as L1 from file A), add all L2s to a new column (pooled-L2), etc.
- In this way, you can have s_{xj} of all stages.

Why are all these difficulties?

- Because L5 and L6 are not the last stage and some N type died in pupal stage, it is therefore not easy to prepare the curve of L6.

You can use similar method that you used for cohort that female and male have different number of stages.



Teşekkür ederim!

سپاسگزارم

謝謝!

Děkuji

Danke!

¡Muchas gracias!

Thank you!

ご清聴ありがとうございます
います!