

read\_me\_first.txt

Gerritsma S, de Haan A, van de Zande L & Wertheim B (2013). Natural variation in differentiated hemocytes is related to parasitoid resistance in *Drosophila melanogaster*. *J. Insect Physiol.* 59:148-158, DOI:10.1016/j.jinsphys.2012.09.017

This folder contains five subfolders with following information:

Description.txt: full citation of the publication

1.Final\_version

PdF of the online version of the publication

2.Manuscript

-Manuscript\_text: Manuscript text

-Tables&Figures

3.Raw\_data

-Dissections\_ER\_22022012.xls: contains the raw data results from dissection experiment of Feb 2012 (Individual encapsulation ability)

-Raw\_data\_Hemocytes\_feb2012.xls: raw data of hemocyte counts feb 2012

-Encapsulation\_Assay\_2010\_sheets\_A\_to\_I.xls: Raw data of encapsulation assay 2010 (Population level encapsulation rate). NOTE: only sheets A to I hold the raw data

-Folder Scanned\_datasheets: contains all raw data from datasheets (scanned) for hemocyte experiments and encapsulation assay (population level encapsulation rate)

4.Processed\_data

-EncapsulationAssay\_2010:

Results encaps essay.xls

R script and datafile for R: EncapsR.txt

-HemocyteCounts&ER\_Feb\_2012

All\_data\_feb2012.xls: holds all the data for both hemocytes and the dissection experiment

R\_Hemocytes\_Data.xls: all the data (hemocytes and ER) in one compiled data file. Can be used for R

R\_Hemocytes\_Data.txt: text file version of previous

Script\_final: R script to analyse the data

Variables\_explanation.doc: explains all the variables used in the R script

-HemocyteCounts\_June\_2011

GOTH\_GRO\_BAY\_KAL\_HemocyteCounts\_June2011.xls: datafile with results from an earlier experiment in June 2011. Data of GOTH was used in discussion

To analyse the data use: GOTH\_GRO\_BAY\_KAL.txt and Script GOTH\_GRO\_BAY\_KAL

-HemocyteCounts\_Okt\_2011

All\_data\_lamellocyte\_exp\_Okt\_Sept\_2011.xls: datafile with results from an earlier experiment in Okt 2011. Only control larvae were tested. Data of GOTH was used in discussion

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To analyse the data use: R\_Lamelloocytes.txt and Script lamelloocyte check

## 5. Protocols

-Dissection&HemocyteCounts: A pdf file describing the whole experimental setup/protocol: Dissection\_Hemocyte\_counts\_protocol.pdf

-EncapsulationAssay\_2010

Screening field lines for encapsulation ability.doc: experimental setup

Schedule encaps assay.xls: time schedule for experiment

## 6. Fieldwork

-Veldwerk\_Samplesheet.xls: contains all information about the captured flies from the natural populations in Europe during fieldwork, including location, number of males, females and species

-Culture\_schedule\_fieldlines\_1stmonths.xls: shows the culture schedule of the fieldlines for the first months in the lab. It also shows when certain flies were merged into one line and when they were at 15 degrees and when at 20 degrees celsius

-Fieldlines\_location.ppt: ppt with google map pictures showing the location of the fieldlines and some extra info

## 7. Scanned labbooks

-Labbook\_AdH\_Hemocyte\_counting\_and\_more: some notes from Ammerins de Haan (second author)

-Pilot\_experiments\_2011: First pilot experiment hemocyte counts

-Labbook1\_SG\_EA\_2010&HC\_2011\_part1: Contains notes on first hemocyte count experiments from 2011 (June/July)

-Labbook1\_SG\_EA\_2010&HC\_2011\_part2

-Labbook2\_SG\_HC\_2011\_2012: Contains notes on hemocyte experiments from Oct/Nov 2011 and the big experiment in Jan/Feb 2012