Annalist

"Data management for the little guys"

Graham Klyne



Drosophila testes gene expression (Fly-TED)









Browse it

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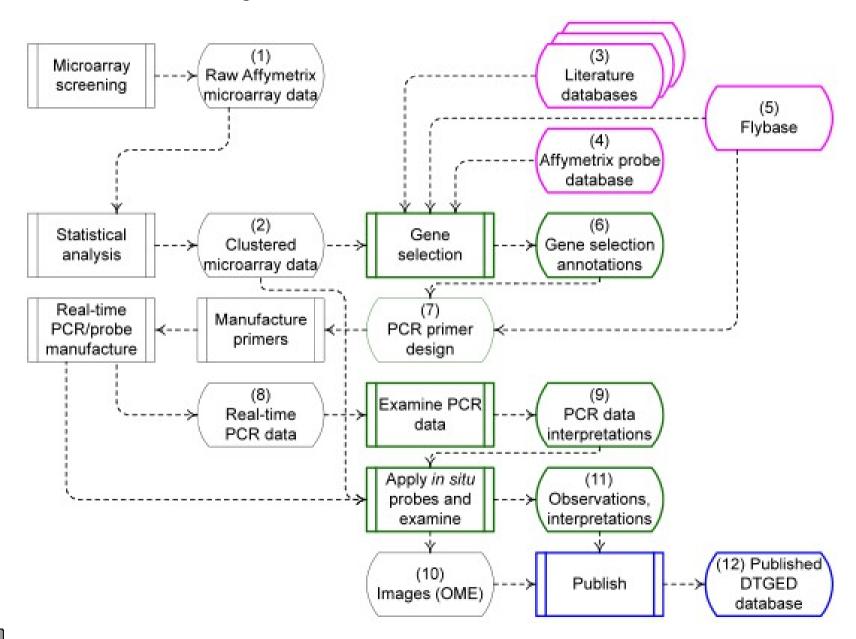
Browse it

- Browse by Gene Name
- Browse by CG Number
- Browse by Strain
- Browse by Expression Location Browse it

http://flyted.zoo.ox.ac.uk/

Excerpts from Fly-TED web site by Jun Zhao Licensed under a Creative Commons Attribution-ShareAlike 3.0 Unported License.

Fly-TED data flows





Fly-TED data management

 Microarray screening and selection of genes microarray data

data collected in spreadsheets

- annotation with literature references

 Real-time PCR and in situ probe manufacture

- more spreadsheets
- In-situ imaging
 - create and examine images of in-situ hybridization



selection

(7)

xamine PCF

Apply in situ probes and

examine

Images (OME)

Clustered

Real-time



Flybase

12) Published



Images: Helen White-Cooper

Literature

Affymetrix probe database

Gene selection

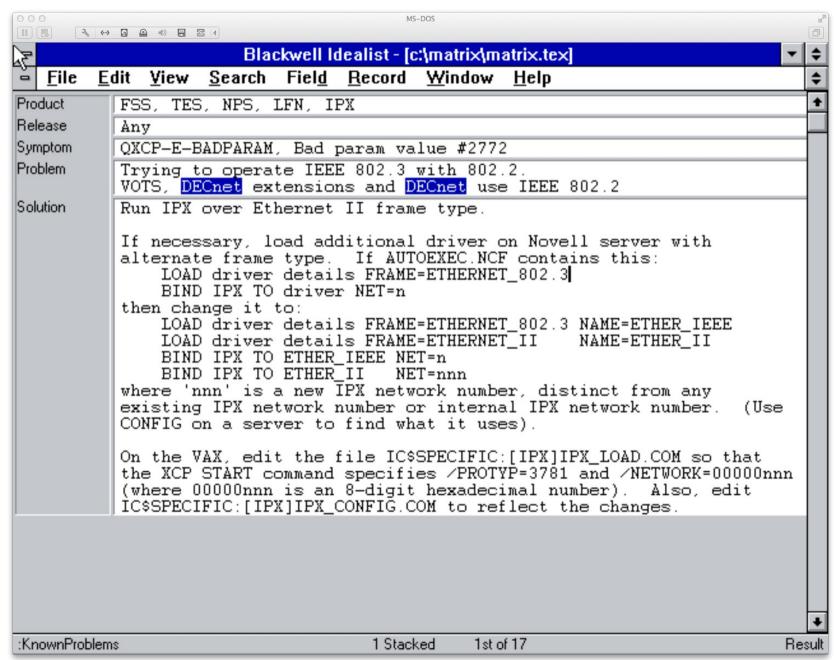
annotations

PCR data interpretations

Observations interpretations

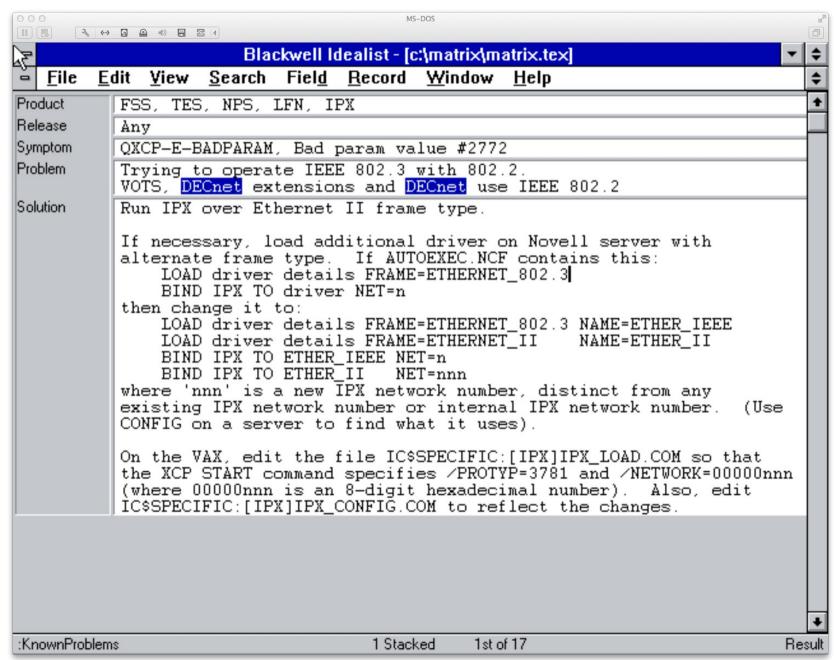
Publish

Technical support knowledge base



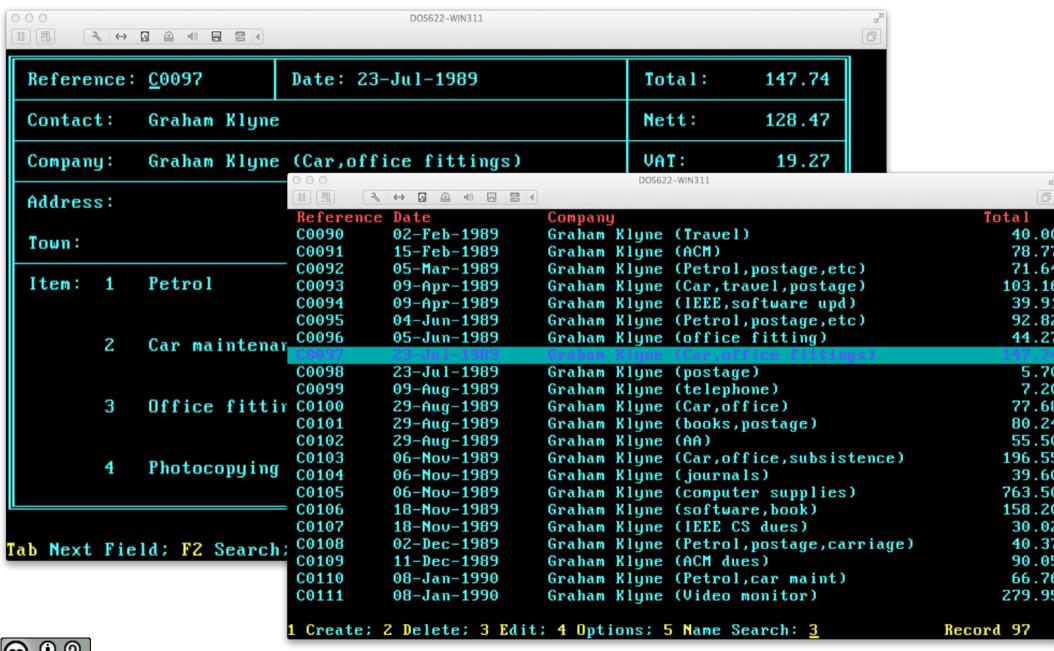


Technical support knowledge base





Small business transactions



Some common themes

Read/write:

- a need for a system that handles both data access and input in the same activity; more notebook than database
- Loosely structured data
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- Structure can evolve as the system is used
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Annalist: mockup demo

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- This demo is a view of what such a system might look like...
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- Usable "out of box"
 - no complicated setup before use ...
 - ... but also a basis for custom applications
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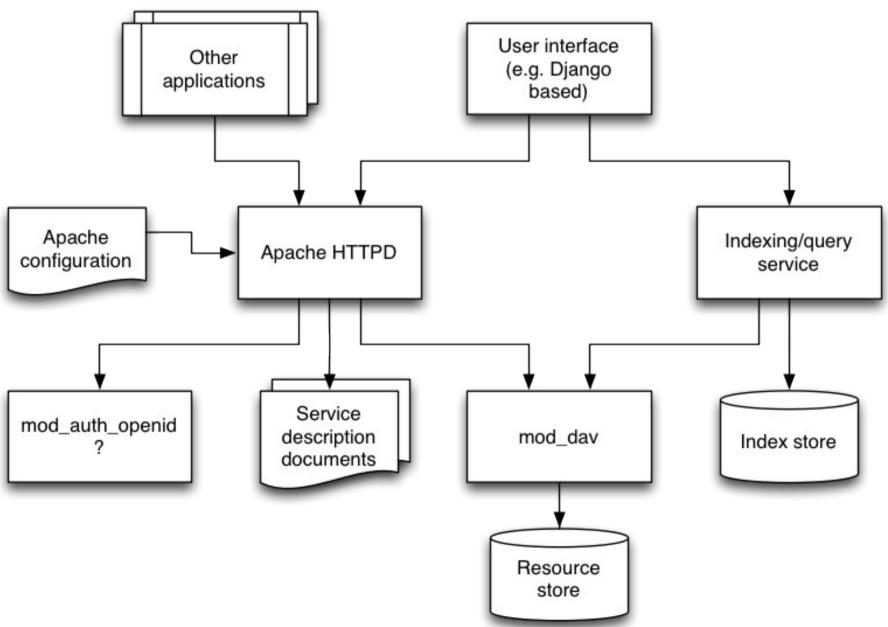


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- Small businesses (SOHO and up...)
- Clubs and non-commercial associations
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- ... where use of data may pass to others who are familiar with different tools



Tentative architecture / overview





Application examples

- Technical support knowledge base
- Small-business bookkeeping
- Personal information management
- 3D model sharing (e.g. Thingiverse)
- Recipe sharing
- Project management

- BioImage / Fly-TED
- Workflow sharing (e.g. myExperiment)
- VIDAAS option (http://vidaas.oucs.ox.ac.uk/)
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 - a commonly requested feature
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I propose to develop and implement a prototype of this design as an independent open-source project in early 2014 (e.g. 50% time over 3-6 months for an initial prototype)

I would like to work on a consulting basis with interested university research activities to develop and refine specific requirements and features (e.g. 25% time)





(A summary slide from 2005)

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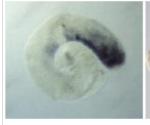
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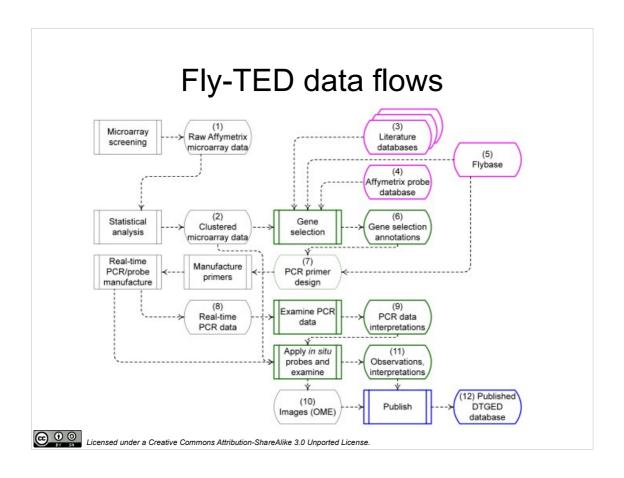
http://flyted.zoo.ox.ac.uk/

Excerpts from Fly-TED web site by Jun Zhao

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One of the first projects I worked on after joining David Shotton's IBRG at Oxford was collection and cataloging of gene expression images of sperm development in Drosophila Melanogaster (a commonly studied model organism), which were collected as part of a strand of research investigating causes of male infertility in humans and other species.

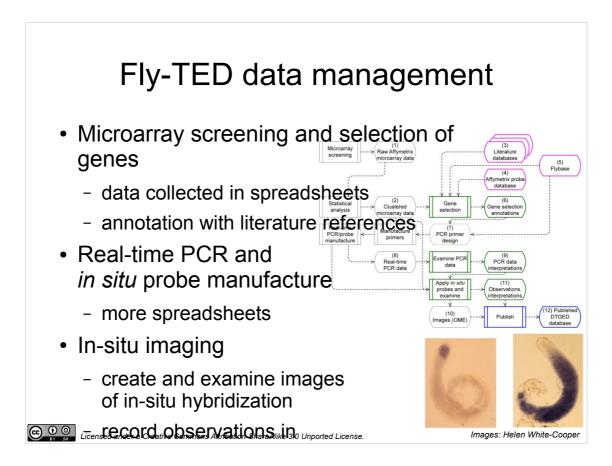
The eventual result of this work was a public database and web site, through which in situ gene expression images can be viewed based on selection by genere name, Drosophila species, expression pattern or other criteria.



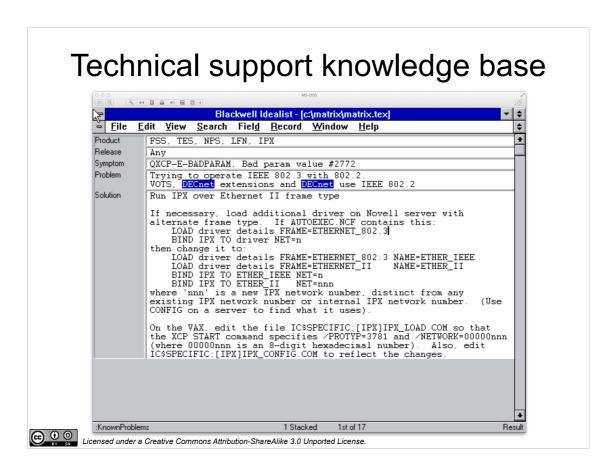
A substantial part of the Drosphila genome was screened, using microarrays, and statistically analyzed to identify genes with distinctive expression patterns in drosophila testes across genetic variants exhibiting male sterility, compared with expression in the normal wild-type variant (e.g. genes in sterile mutants whose expression was significantly higher or lower than in the wildtype).

This screening was used in combination with informed researcher evaluation to select a subset of about 1000 genes for in-situ gene expression imaging, a complex and labour-intensive process that results in an image showing where in the sperm development cycle the various genes are expressed.

This work required the collation and collation of diverse information about a range of drosophila genes associated with the microarray "probe" sequences: gene identification, species variant identification, microarray expression levels, statistical analyses, real-time PCR data, in-situ images, image annotations (including sperm development stage) based on manual through-microscope observation of the images.



For this work, data on the genes selected for in-situ imaging was collected in spreadsheets with file-system references to the corresponding in-situ images. Because the spreadsheets were manually populated by a number of different people, some of the image annotations, such as those of the sperm development stages, were not consistently described and needed manual review and alignment when the data were prepared for publication

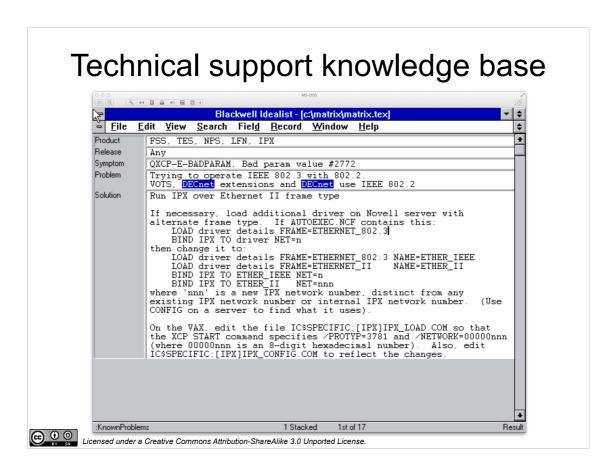


Many years ago, in the early days of personal computer adoption in corporate environments, I helped support a range of software products for connecting diverse kinds of computers, specifically between PCs and mainframe computers.

We were providing person-to-person technical and development support to high-level technical staff in large corporations (banks, telecoms, legal, multinational consumer product companies, etc.), so good access to product information, common failure modes and remediation was important. Things one might find today in a Google search returning a page on Stack Exchange, but before the Web was widely used.

I used an early PC software package called "Idealist" to create a knowledge base that could be searched online by any or all of keyword, product details, operating system version, etc. to locate possible causes and remedies for a reported problem.

Idealist was unusual in that is was one of the first PC software packages to provide free-text keyword searching in semi-structured data. When a new problem was encountered, a description and summary would be entered into the knowledge base by a technical support person so that information about it would be available should it occur for any other customer reporting the same problem. Idealist was probably ahead of its time, as it did not survive as a commercial product, and as far as I'm aware there is still not a ready-to-go product that provides the same level of lightly structured data entry and structured search. (There are plenty of developer tools that might be used to create such a product, but not one I'm aware of that just works out of the box.)



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For many years, I ran a small self-employed consulting business, starting in the early days of availability of personal computers at a price that individuals and small businesses could afford. I had a need to raise invoices, handle purchase orders, etc.

Not wishing to pay for a fully-fledged accounting software package, I used a combination of WordPerfect and an associated utility called Notebook to create a kind of computerized card index, with a "card" for each invice, purchase order, etc. The Notebook utility maintained its data in a simple text document format that was compatible with WordPerfect's mail-merge capabilities, so could easily be formatted as a document for submission to a customer, supplier or just filed.

The simple, out-of-box working of Notebook, and its integration with other WordPerfect office suite programs, made it very easy for me to set up a system that maintained records and generated appropriate presentations, which I used for many years. (I now use a descendant of that system based on MS Access, which was much harder to set up and is, in many ways, less satisfactory in use.)

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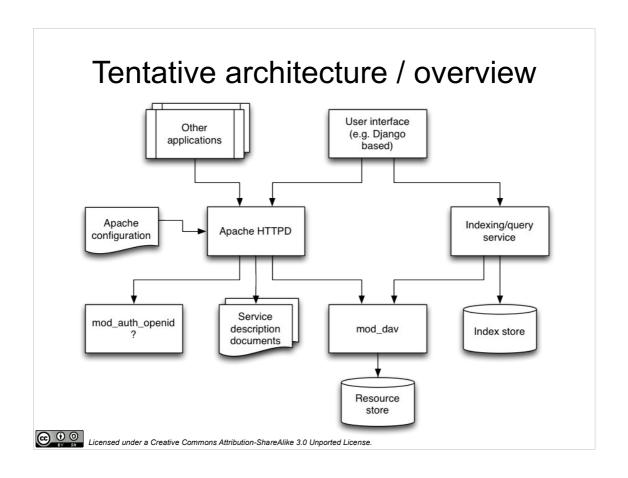


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