



***B*IODIVERSITY INFORMATION
RETRIEVAL ACROSS NETWORKED
DATA SETS**

By

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Sir Robert May (1998)

*“... The next century will be the ‘**A**ge of Biology’, just as [the twentieth century was] the age of physics and astronomy. Specifically, those countries who best know how to correlate, analyze and communicate biological information will be in the leading position to achieve economic and scientific advances.”*



*B*iodiversity information:

disperse
unreported
inaccessible



BIODIVERSITY INFORMATION: MALAYSIAN PERSPECTIVE



Identified as one of the world's twelve mega-diversity areas with extreme rich biological resources

*over 15,000 species of higher plants,
300 species of mammals,
254 species of breeding birds,
198 species of amphibians,
379 species of reptiles,
over 150,000 species of the invertebrates,
over 4,000 species of marine fishes
449 species of freshwater fishes in Malaysia*

(Earth Trends, 2003)

Current status

Some institutions have information documented in the form of **databases**, while other institutions are looking into it

Databases are in **heterogeneous** formats.

Few databases are on the Web, while many are available **offline**

Some of these databases are well-structured, others are largely project /species specific and/or **unstructured**

These databases exist **independently**

There is no **meta-data** (data- dictionary)



NO FRAMEWORK TO LINK THE
SCATTERED DATA SO AS TO
FACILITATE EXCHANGE OF DATA
AMONGST THE DIFFERENT DATABASES
AVAILABLE IN THE COUNTRY.

Need to overcome this setback....

HOW??

*Dissemination and retrieval of
information in a networked
environment....*

COMMUNICATION of biodiversity
information (*biodiversity informatics*)

LINK all the disperse information
clusters in remote and distributed
medium



THE SOLUTION





Rationale for indigenous DATABASE INTEGRATION SYSTEM

Implementation issues existing
systems
No control



INDIGENOUS framework for integrating **DISTRIBUTED**
HETEROGENEOUS *relational*
biodiversity databases

to facilitate biodiversity data sharing among the scientific
contemporaries especially in Malaysia.

Philosophy:

Database **KEPT & MAINTAINED** by DATA OWNER

SIMPLICITY is stressed to offer a **USABLE** system for the
scientific community in Malaysia and future integration with
other global networks.



The FOCUS

D DEVELOPMENT OF A NEW DATABASE INTEGRATION
SYSTEM

D DEVELOPMENT OF BIODIVERSITY DATABASES
ACCORDING TO REQUIREMENTS SET BY PARTNERS



***RELATIONAL
BIODIVERSITY
DATABASES***



DATABASE INTEGRATION



*P*ROPOSED DATABASE INTEGRATION

SYSTEM

- *Links remote databases on networked environment.*
- *Supports heterogeneous data format.*
- *Supports heterogeneous database management systems.*
- *Links databases hosted in Windows and UNIX based platforms.*
- *Provides data security for database owners by allowing them to keep and maintain their own data and to choose information to be shared and linked.*



PROTOTYPE

Microsoft Windows

ASP

PHP

XML

HTML

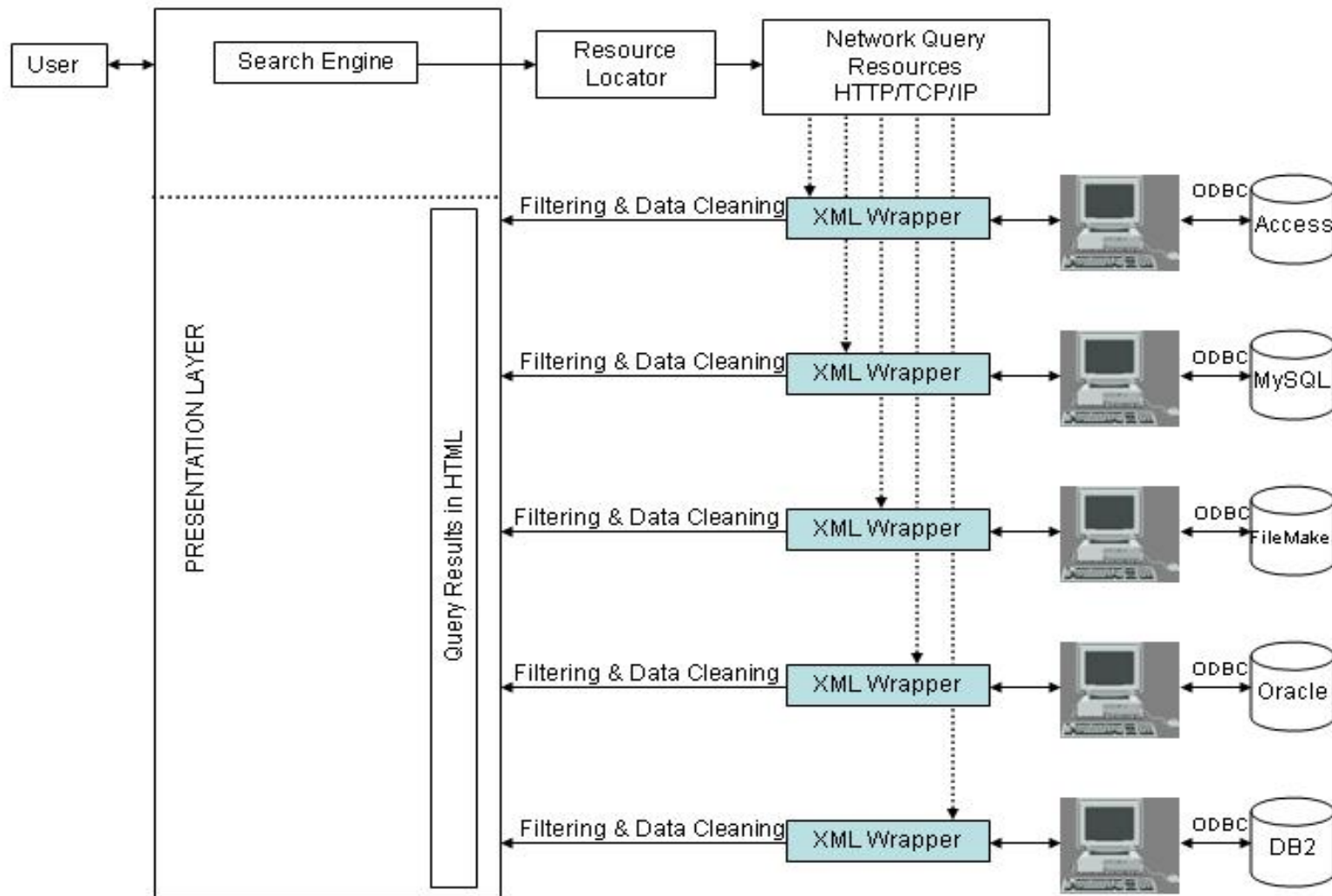
VBScript and Jscript

DBMS- FileMaker, mySQL, DB2..



Search Mechanism

ARCHITECTURE



SUMMARY

- a. Created data format & relational biodatabases
- b. Developed indigenous integration system

- c. In summary

Achieved objectives set out which are:-

- *Links remote databases on networked environment.*
- *Supports heterogeneous data format.*
- *Supports heterogeneous database management systems.*
- *Links databases hosted in Windows and UNIX based platforms.*
- *Provides data security for database owners by allowing them to keep and maintain their own data and to choose information to be shared and linked.*



FUTURE FOCUS

To further develop & enhance this biodiversity content architecture for our country

- ❖ e-library (Parasitofaunae Malaysiana)
- ❖ Data-security
- ❖ Use of Semantic technology
- ❖ Populate Relational Databases



RESEARCH PROJECTS

Parasites Informatics Network System (PINS)

In our effort to aid research in Parasitology in Malaysia we have developed PINS which is essentially a parasite taxonomic database with inputs from experts in the field who will maintain their respective database with technical assistance from us. We will be incorporating into this database related IT Alternatives to Taxonomy which are currently developing as well as ecological and biological data which will be linked semantically.

[Sarinder Kaur & Lim LHS and other researchers]



Image & text recognitions for taxonomy: retrieving, managing, enhancing and recognizing different types of images

This project is initiated to develop an image-recognition system to aid in species identifications (taxonomy). It is first necessary to develop a database to store images in different formats either from our research or retrieved from different sources. The retrieved images will probably have to be enhanced (if necessary) for image recognition. Our query to search and capture images from publications will be textual in nature since most of the images and illustrations are tagged textually. The system we are developing is for monogenean species, a parasitic flatworm.

[Arpah (PhD candidate), Sarinder Kaur & Lim LHS]



2-D to 3-D modeling of organisms for structural morphology

Visualisation in 3-D provides more information on the spatial relationships of objects or structural morphological features in the case of organisms. 3-D visualization is important in understanding structural and functional morphology and also important for taxonomic and identification purposes. The aim of this study is to construct 3-D models from 2-D images. In this study several softwares capable of doing this have been evaluated and we are currently using 3-d Max programme because of its advantage of being able of rendering irregular shapes using polygons.

[Teo Bee Guan (MSc candidate), Lim LHS, Sarinder Kaur]



Image recognition using DAISY

This project looks into using an available software on image recognition for taxonomy to aid in identifications. The group of interest is to use the software to image moths which have different shapes, sizes and colour patterns. This project involves setting up a database of selected groups of moths.

[Evelyn (MSc candidate), Sarinder Kaur & Lim LHS]



THANK YOU

