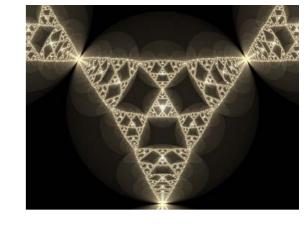


Teratology Society, Monterey CA, July 1, 2008

Thomas B. Knudsen, PhD
National Center for Computational Toxicology
US EPA, Research Triangle Park NC
Knudsen.Thomas@epa.gov

DISCLAIMER: views reflect those of the presenter and not EPA policy; any reference to commercial products or tradenames does not constitute endorsement

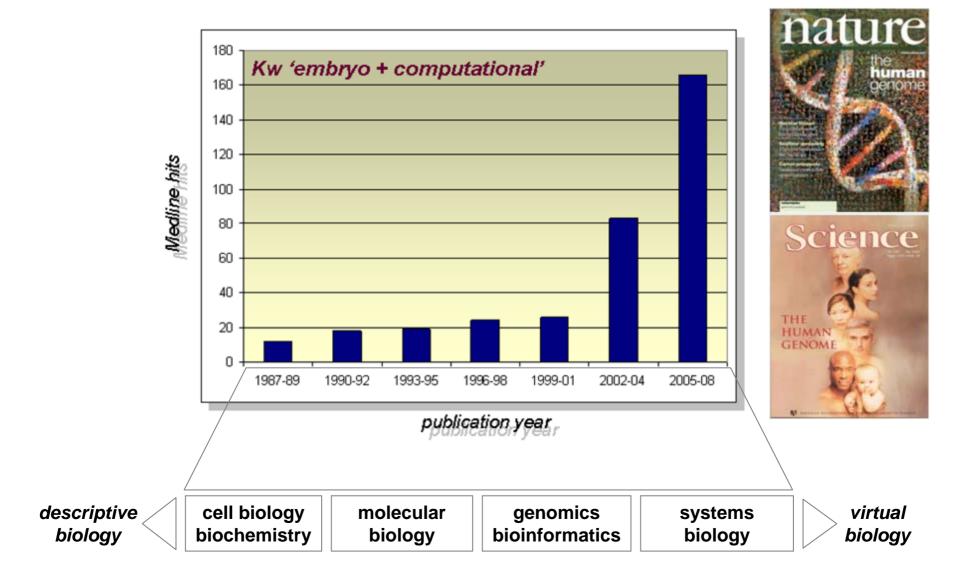
INTRODUCTION



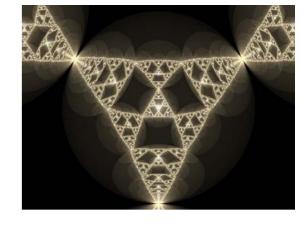
- computers, imaging, and the worldwide web have assumed an important role augmenting learning
- since mid-1990's web resources have become available to accelerate research and advance knowledge
- technologies to help computers access scientific information fuel drive to make it 'freely available to all'
- efforts needed to prevent a 'free for all' when elevating these data into computable knowledge

Computational (in silico) biology:

impact of the human genome project



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Virtual Libraries



About SDB Membership

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Publications

Virtual Library

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Virtual Library-Developmental **Biology**



This Virtual Library maintained by Society for Developmental

Biology.

DEVELOPMENTAL

Published by Elsevier Science under Auspices of Society for Developmental Biology

Subject Index

Gametogenesis and Fertilization

Early Development

Organogenesis and

Morphogenesis

Pattern Formation Gene Regulation and

Genetics

Cell Lineage and Fate

Maps

Evolution and Development

Diseases, Defects and

Development

Research Resources Index

Departments and Institutes Index

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Search Engines and Other

Societies and Organizations Resources

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Vertebrates Invertebrates

Plants |

Unicellular/Lower Eukarvotes and

Prokaryotes

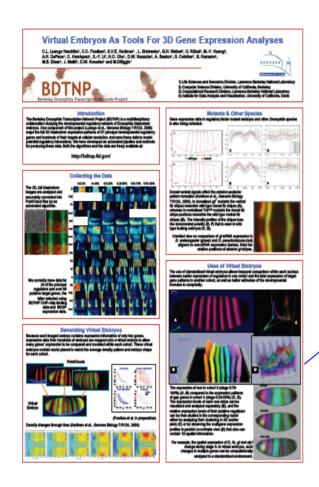
Journals and Publishers

Index

Journals Publishers

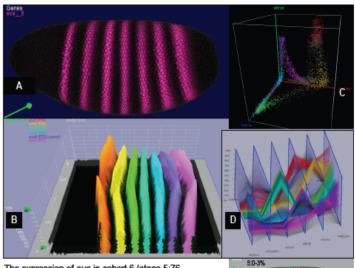
Educational Resources

- Stephen Alexander's laboratory at the University of Missouri uses Dictviostelim discoideum as a model system for studies on: 1, molecular and cellular mechanisms of resistance to anticancer drugs: 2. regulation of protein secretion during development.
- Zena Werb's lab at UC San Francisco studies the role of extracellular proteolysis in controlling vascular development and angiogenesis during embryonic development and placental formation, bone development and tumorgenesis.
- Tumor Suppressor Genes in Drosophila and Their Human Homologs; Genetics of Development in Drosophila The objective of research in Peter J. Bryant's laboratory (UC Irvine) is to understand how cell proliferation is controlled during development, and how genetic mutations lead to growth abnormalities and cancer. They approach this problem using a genetic approach in Drosophila, by identifying and characterizing tumor suppressor genes, in which mutations cause excessive cell proliferation in the developing organs of mutant larvae. Efforts from this lab and others have led to the identification of over 60 of these genes, and over 20 of them have been cloned. Almost all of them have human homologs whose potential roles in cancer are now being investigated.
- Atlas of Developmental Abnormalities in Common Laboratory Mammals This website is designed for rapid access to the image(s) of developmental abnormalities in common laboratory mammals.



Uses of Virtual Embryos

The use of standardized virtual embryos allows temporal comparison within each nucleus between earlier expression of regulators in one cohort and the later expression of target gene patterns in another cohort, as well as better estimates of the developmental increase in complexity.



The expression of eve in cohort 6 (stage 5:76-100%) (A, B) compared to the expression patterns of gap genes in cohort 4 (stage 5:26-50%) (C, D). The expression levels of each eve stripe can be visualized and analyzed separately (B), and the relative expression levels of their putative regulators can be then studied in the corresponding nuclei either by analyzing their clustering in 3D scatter plots (C) or by observing the multigene expression profiles in parallel co-ordinate view (D) that also can contain 1D spatial information.

For example, the spatial expression of *D*, *Kr*, *gt* and *slp1* change during stage 5. In virtual embryos, such changes in multiple genes can be computationally analyzed in a standardized environment.

551-75%

Source: Berkeley Drosophila Transcription Network Project (BDTNP)

http://www-vis.lbl.gov/Publications/2007/LBNL-63062.pdf

Virtual Library needed for teratology

- relaunching of Society's website coming soon!



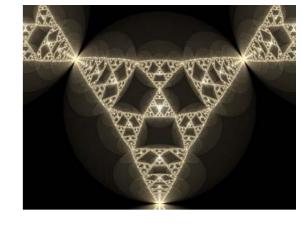
Fair Use Doctrine in education:

http://www.dartmouth.edu/~webteach/articles/copyright.html

- purpose: not-for-profit use 'fair' if purpose of web content is instructional
- nature: works created for research purposes likely 'fair' due to expectations for educational use
- substantiality: 'fair' use if reasonable relative to the whole, or if not the basis of user's work
- 4. marketing: 'fair' if use has little or no effect on creator's ability to market his/her work

<u>BEST PRACTICE</u>: assume work copyrighted when web page created; get author's consent and acknowledge; disclose that materials fall under fair use exemption and are restricted from further use

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Digital atlases: Carnegie Human Embryology Collection

The Virtual Human Embryo

http://nmhm.washingtondc.museum/collections/hdac/index.ht

image databases for 23 stages of human embryonic development; each includes a complete serially sectioned embryo from the Carnegie collection

Multi-Dimensional Human Embryo

http://embryo.soad.umich.edu/

web-accessible 3D image reference resource based on magnetic resonance microscopy of the Carnegie Collection

Multi-Dimensional Human Embryo

Sample images listed by: Carnegie Stages



Home

Atlas

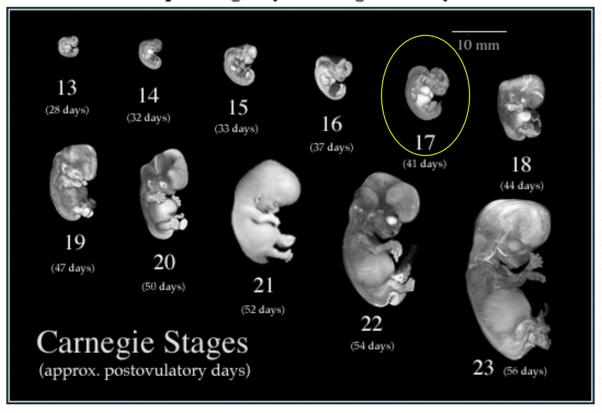
Project

Technical

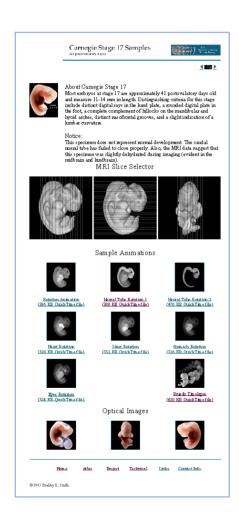
Links

Contact Info

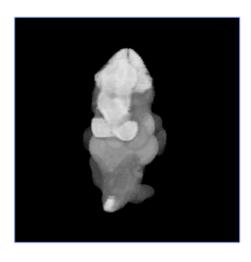
View Sample Images by Selecting an Embryo Below

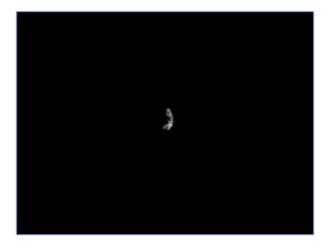


Carnegie Stage 17 Samples









Online textbooks

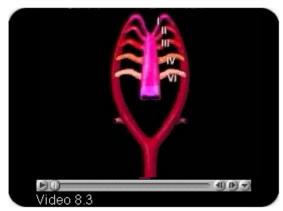


Release 1.0 @ 2001 Kyle E. Rarey, PhD and Lynn J. Romrell, PhD

Source: http://www.imc.gsm.com/siteinfo/browse_heinfo.htm

Chapter 8: Heart Development, discussion of aortic arches





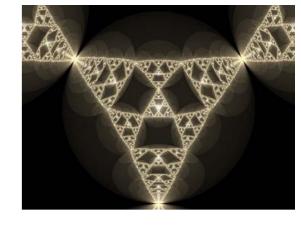
Arterial Outflow Tract

The arterial outflow tract of the primitive heart is composed of the aortic sac and aortic arch arteries. Blood from the truncus arteriosus passes into the aortic sac and then into bilateral pair of six aortic arches that are located within the pharyngeal arches. Blood then enters paired dorsal aortas that unite to become one medial dorsal aorta below the heart. Branches from the dorsal aorta are vitelline arteries, umbilical arteries, and intersegmental arteries.

The truncus ateriosus eventually becomes part of the ascending aorta and the proximal portions of the pulmonary arteries. The aortic sac forms part of the aortic arch and part of the brachiocephalic artery. The adult derivatives of the six, paired aortic arches are listed in the following table (video 8.3 - developing arterial outflow seen from an anterior view).

Aortic Arch Artery	Adult Derivative			
1st	Maxillary artery			
2nd	Stapedial artery, hyoid artery			
3rd	Common carotid arteries, roots of internal carotid arteries			
4th	Part of the arch of the aorta, right subclavian artery			
6th	Ductus arteriosus (left side), roots of pulmonary arteries			

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Tour the new MGI website

ID or symbol or name

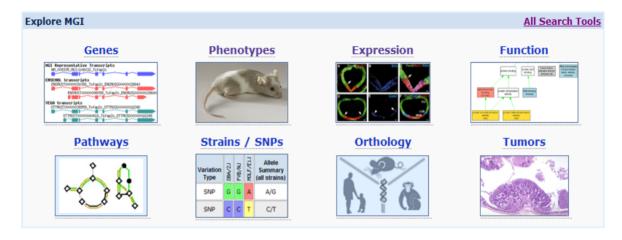
Ouick Search

Mouse Genome Informatics

http://www.informatics.jax.org/

Search ▼

Download ▼ More Resources ▼ Submit Data Find Mice (IMSR)



FAO₅

How do I...

- .. search for genes? FAQ
- .. find mutations for phenotypes or diseases? FAQ
- .. find expression data? FAQ
- .. view a structural genomic map? FAO

More FAQs

News

17 June, 2008

- · MGI genome coordinates are now updated to NCBI Mouse Build 37 and dbSNP Build 128.
- The MGI website and navigation tools have been redesigned. Take a guick tour.
- MouseCvc, a new database of curated biochemical pathways, is now available at MGI.

More MGI news

MGI Statistics

Mouse Community News

MGI Roadshow: If you would like to host a MGI workshop at your institution, contact User Support to discuss the details.

Visit MGI posters at these upcoming conferences:

- XX International Congress of Genetics
- International Conference on Bioinformatics & Computational Biology
- International Conference on Intelligent Systems for Molecular Biology

Read about the International Mouse Knockout Consortium [KOMP (USA); EUCOMM (Europe); NorCOMM (Canada); TIGM (USA)] in Cell 2007; 128: 9-13, PMID: 17218247 and Cell 2007; 129: 235, PMID: 17448981.

7

MP term:

Synonym:

Search ▼

abnormal eye development

MP id: MP:0001286

Definition: malformation or arrest of differentiation of the visu

Number of paths to term: 1

denotes an 'is-a' relationship denotes a 'part-of' relationship

Phenotype Ontology

vision/eye phenotype

abnormal eye morphology

abnormal anterior eye segment morphology +

■ <u>abnormal eye development [MP:0001286]</u> (278 genotypes, 38)

abnormal corneal stroma development

abnormal eye muscle development

abnormal lens development +

abnormal optic cup morphology

abnormal optic eminence morphology

■ abnormal optic stalk morphology

■ abnormal optic vesicle formation +

abnormal periocular mesenchyme morphology

abnormal retinal progenitor morphology

I aniridia

■ coloboma

cvclopia

ectopia lentis

Illenticonus

I persistence of hyaloid capillary system

■abnormal eye distance/ position +

■ abnormal eve muscle morphology +

abnormal eye pigmentation +

abnormal eve size +

■ abnormal evelid morphology +

■ abnormal lacrimal gland morphology +

abnormal orbit morphology +

abnormal posterior eve segment morphology +

abnormal sclera morphology +

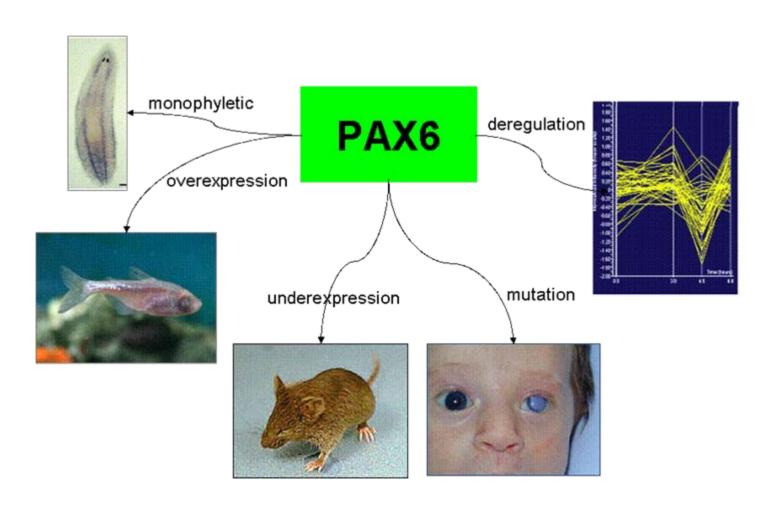
abnormal uvea morphology +

■ intraocular calcification +



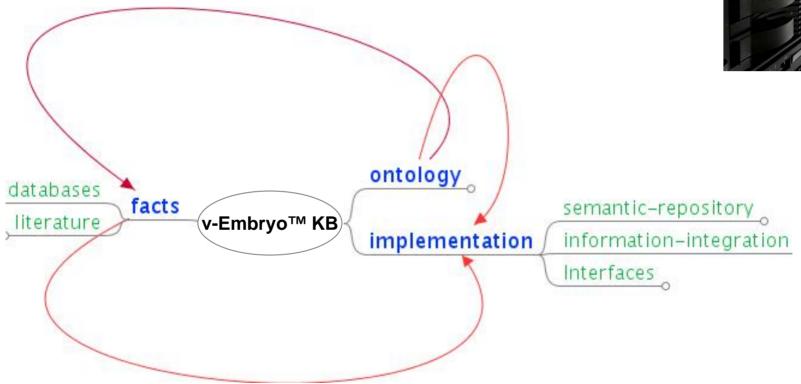
4 genotypes with 4 annotations displayed						
Searched Term: aniridia						
Allelic Composition (Genetic Background)	Annotated Term					
Mab21l1 ^{tm1Nac} /Mab21l1 ^{tm1Nac} (involves: C57BL/6)	aniridia	<u>J:82312</u>				
Pax6 ^{Sey-Dey} /Pax6 ⁺ (C3H/He)-Pax6 ^{Sey-Dey})	aniridia	<u>J:10820</u>				
Pax6tm1Gfs/Pax6tm1Gfs (involves: 129S7/SvEvBrd)	aniridia	<u>J:76576</u>				
Tcfap2a ^{tm1Will} /Tcfap2a ^{tm1Will} (involves: Black Swiss)	<u>aniridia</u>	<u>J:52402</u>				

Pax6 roles in eye development from analysis of the scientific literature



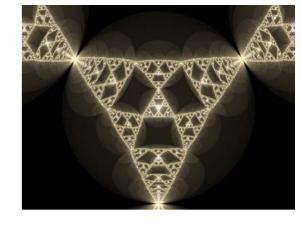
v-Embryo[™] Knowledgebase: normal and abnormal embryogenesis





Source: Imran Shah, NCCT – EPA http://www.epa.gov/ncct/index.html

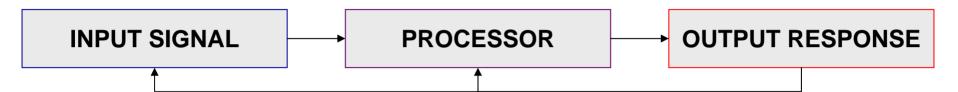
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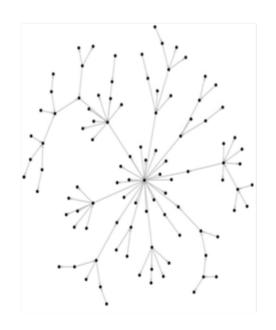
Developmental signaling pathways:

integrating these data at a systems level



Developmental Signals

Wnt, TGFβ, Shh, RTK, Notch-Delta, NF-kB, PCD, nuclear hormone receptors, RPTPs, receptor GC, cytokines, NO, GPCRs, integrins, CADs, gap junction, ligand-gated cation channels, UPR, p53



Morphoregulatory Responses

patterning
proliferation
apoptosis
differentiation
adhesion
motility
shape
ECM remodeling

Edinburgh Mouse Atlas Project



3D digital atlas | theiler stage selector

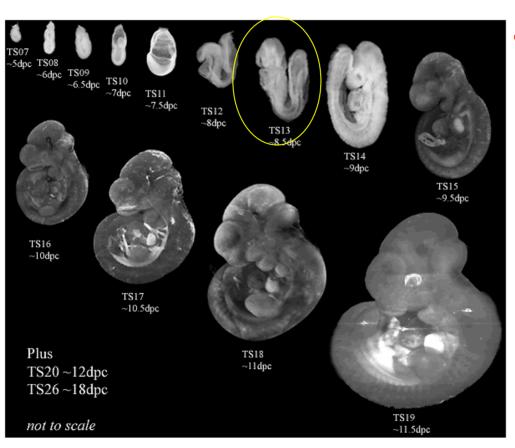
HOME 3D DIGITAL ATLAS

EMAGE DATABASE

RESOURCES

CONTACT

SITE SEARCH



emap models available:

(()

Select the embryo you want

Theiler Staging Guide:

Individual Stages

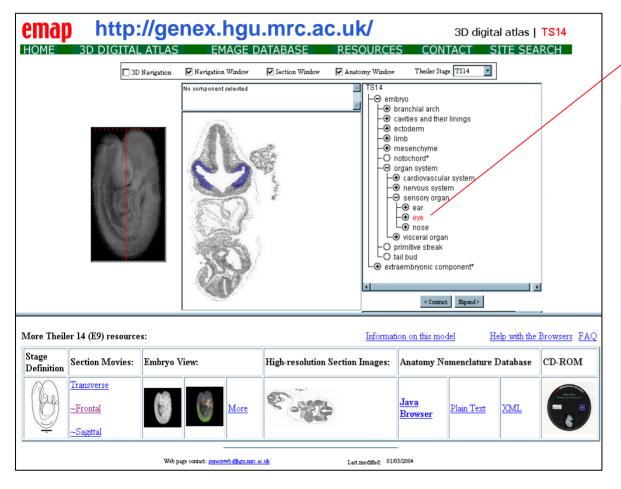
1	2	<u>3</u>		5		7
8	9	<u>10</u>	<u>11</u>	<u>12</u>	<u>13</u>	<u>14</u>
<u>15</u>	<u>16</u>	<u>17</u>	<u>18</u>	<u>19</u>	<u>20</u>	<u>21</u>
<u>22</u>	<u>23</u>	<u>24</u>	<u> 25</u>	<u> 26</u>	<u>27</u>	<u>28</u>

All Stages

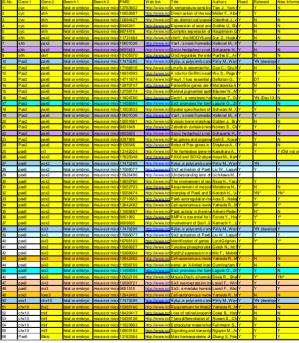
Criteria Summary table		
Pictorial stage index		
Text anatomy index		
Anatomy Database		

FAQs

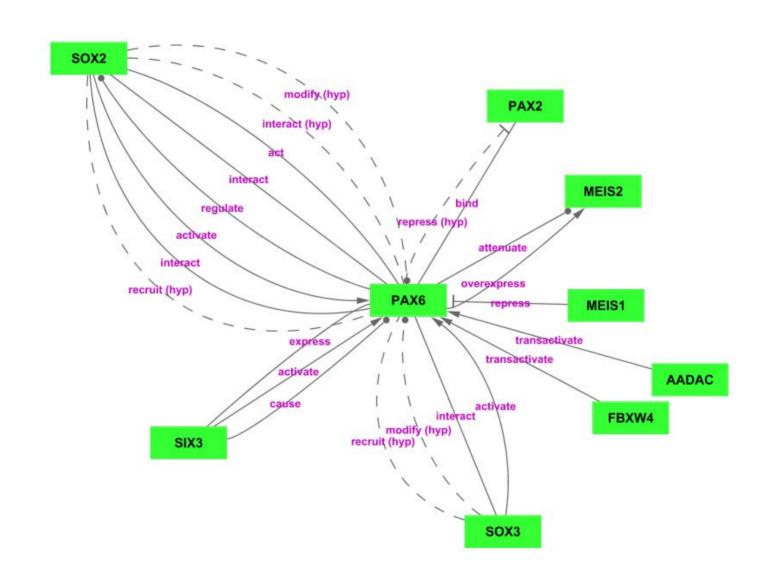
Capturing data from EMAGE, GXD and the scientific literature



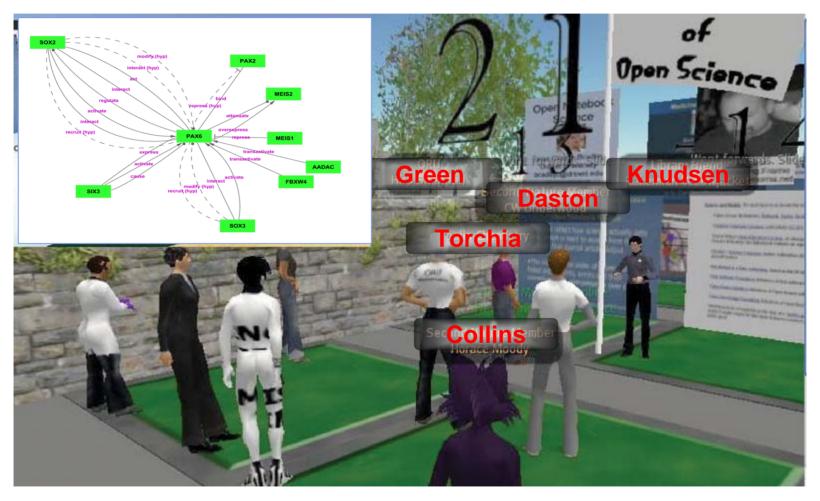
63 genes (TS12-18) captures 2164 PMIDs



EXAMPLE: Pax6 network in eye development

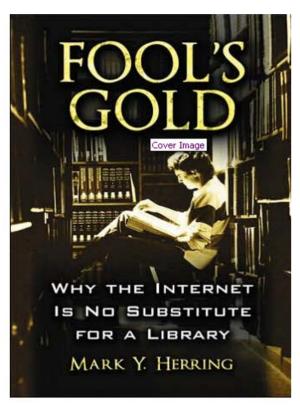


VIRTUAL WORLD: open discussion forum led by Avatars in the virtual-reality world of 'Second Life'



Source: Figure 1, Murray-Rust (2008) Chemistry for everyone. Nature 451: 648-651

Internet extends but does not replace a traditional learning environment



Dacus Library, Winthrop University Macfarland & Company (2007) ISBN-13: 9780786430826

- Herring explores whether the internet will obviate need for traditional print-based academic libraries ... NOT!
- pitfalls & prevarications of popular search engines and digitization of traditional print material
- argues for library's staying power in face of technological advancement