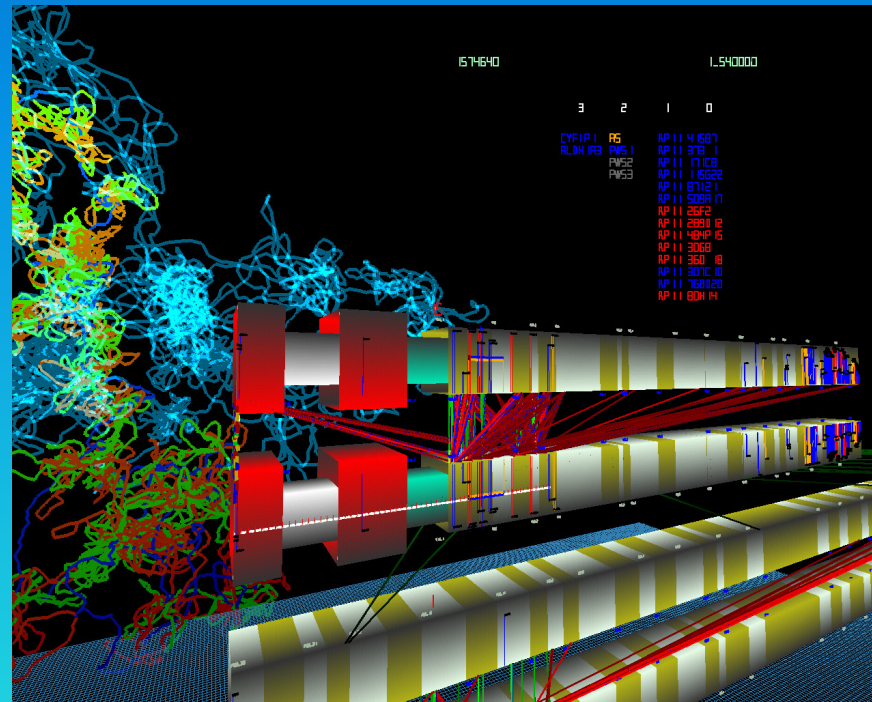


# The GLOBE 3D Genome Browser

GLOBE Genome Information Systems Solutions



to navigate and discover  
the secrets of genomes



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# The Adventure to Discover the Secrets of Life

## Better Health and Longer Living

Genetic information is at the heart of every living being !

### Genome Research, Products & Diagnostics

- extremely huge
- extremely complicated

### >>> The Human Genome

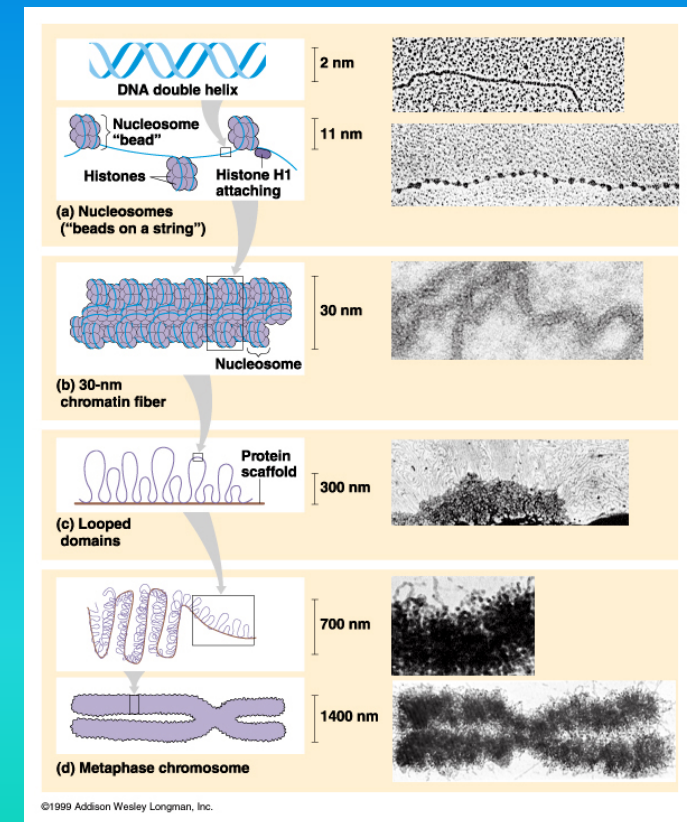
- 3,000,000,000 base pairs
- 30,000 genes
- >1,000,000 disease relevant mutations

### Research & Development

- genome organization and function
- genotype-phenotype correlation
- personal genome projects

### Genomic Products & Diagnostics

- >5,000 genetic tests on the market
- per test often >10,000 genomic markers



>>> e.g. breast cancer, cardiovascular disease, prenatal diagnostics

# The Challenge to Tame the Genomic Beast

## The Requirements/Needs of the Genomic Sector

Adequate Platform to Exploit the Genomic Era !!

### *Generals*

- visualization of data and relations
- combining public and experimental data
- integration from base pair to chromosome

### *Research & Development*

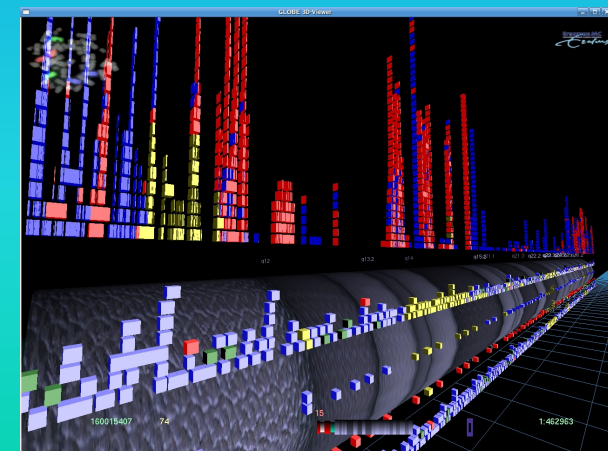
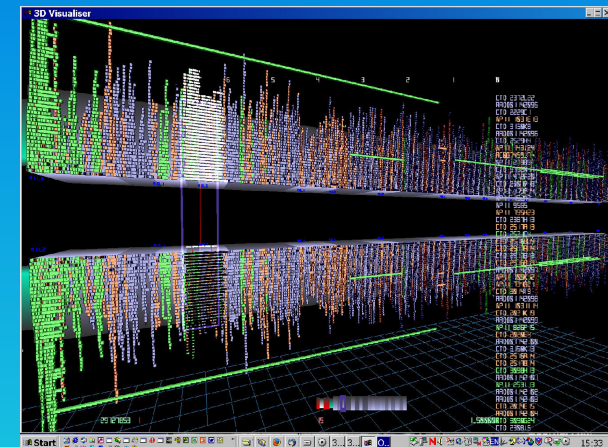
- working at the knowledge frontier
- flexibility, customizable, real-time
- annotation of data and experiment design

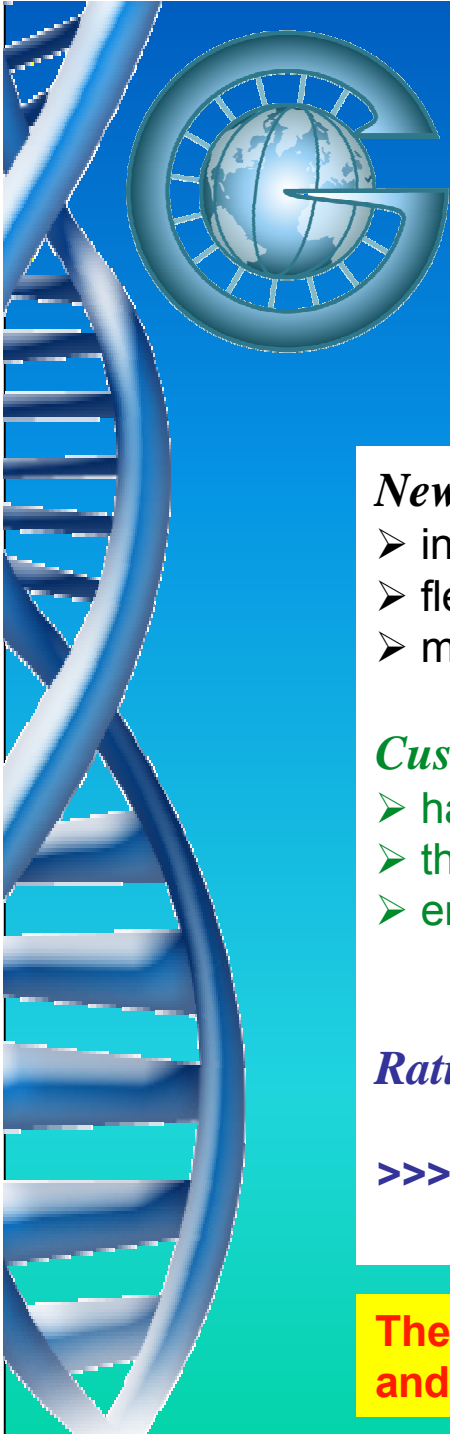
### *Genomic Product Suppliers*

- catalog and sales of complex products
- customizable products on demand
- comparison to other products

### *Diagnostic Institutions*

- easy visual access to diagnostic results
- links to disease data bases for diagnosis





## **Solution**

**Our New Standard for Integrated  
Scientific, Clinical and Industrial  
Genome Research, Application and Exploitation**

### ***New Concept of Data Navigation***

- intuitive and easy to use
- flexible and generic
- multi-dimensional and real-time

### ***Customer Benefits***

- handier and faster to work with
- things which could not be done before
- entire new options for customers

### ***Rationale: A Genomic Global Positioning System (GPS)***

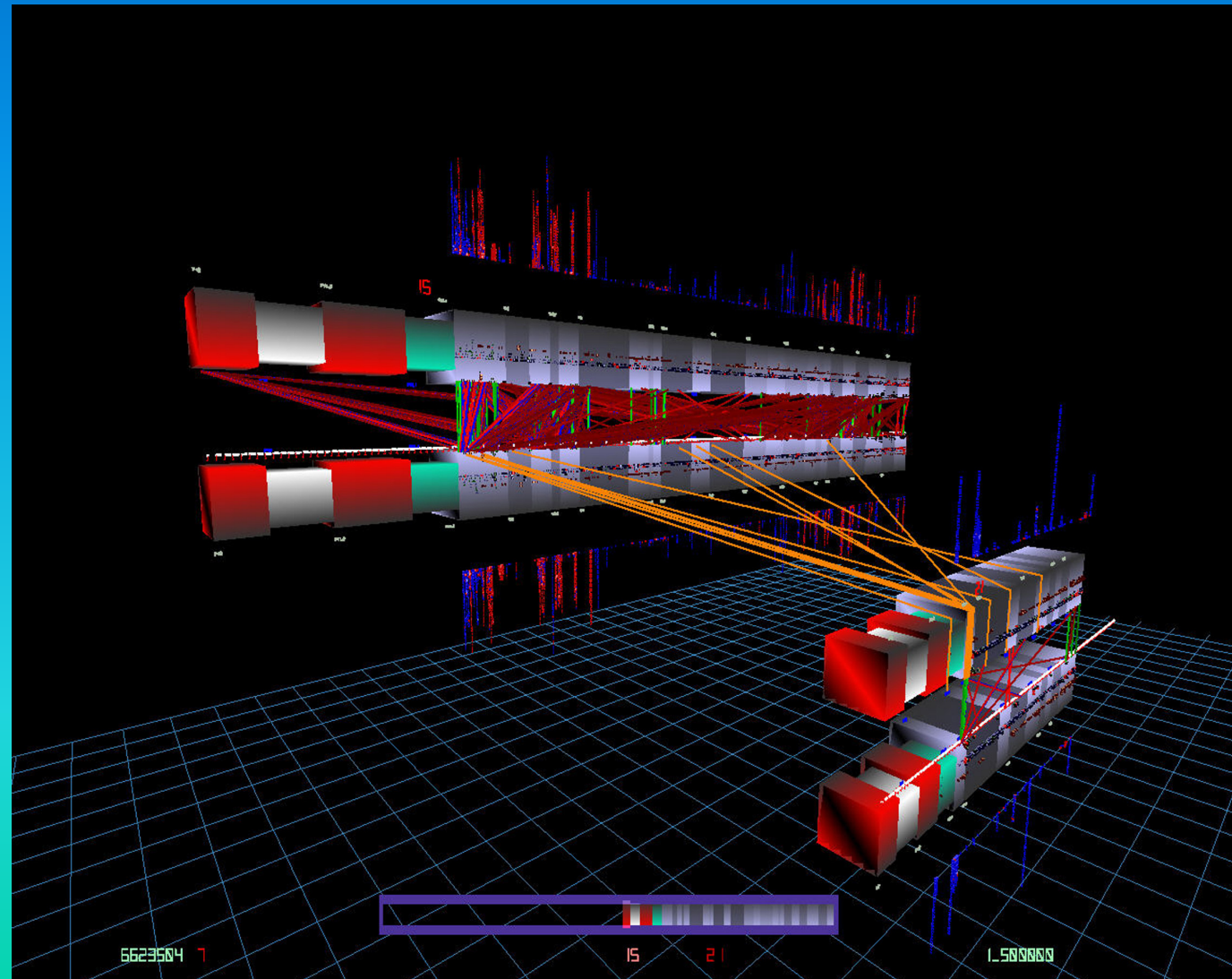
**>>> street list >>> ATLAS >>> Google Earth >>> electronic TOMTOM**

**The GLOBE 3D Genome Browser is the Genome Positioning System  
and Google Earth / Excel / Photoshop in a Single Platform!**



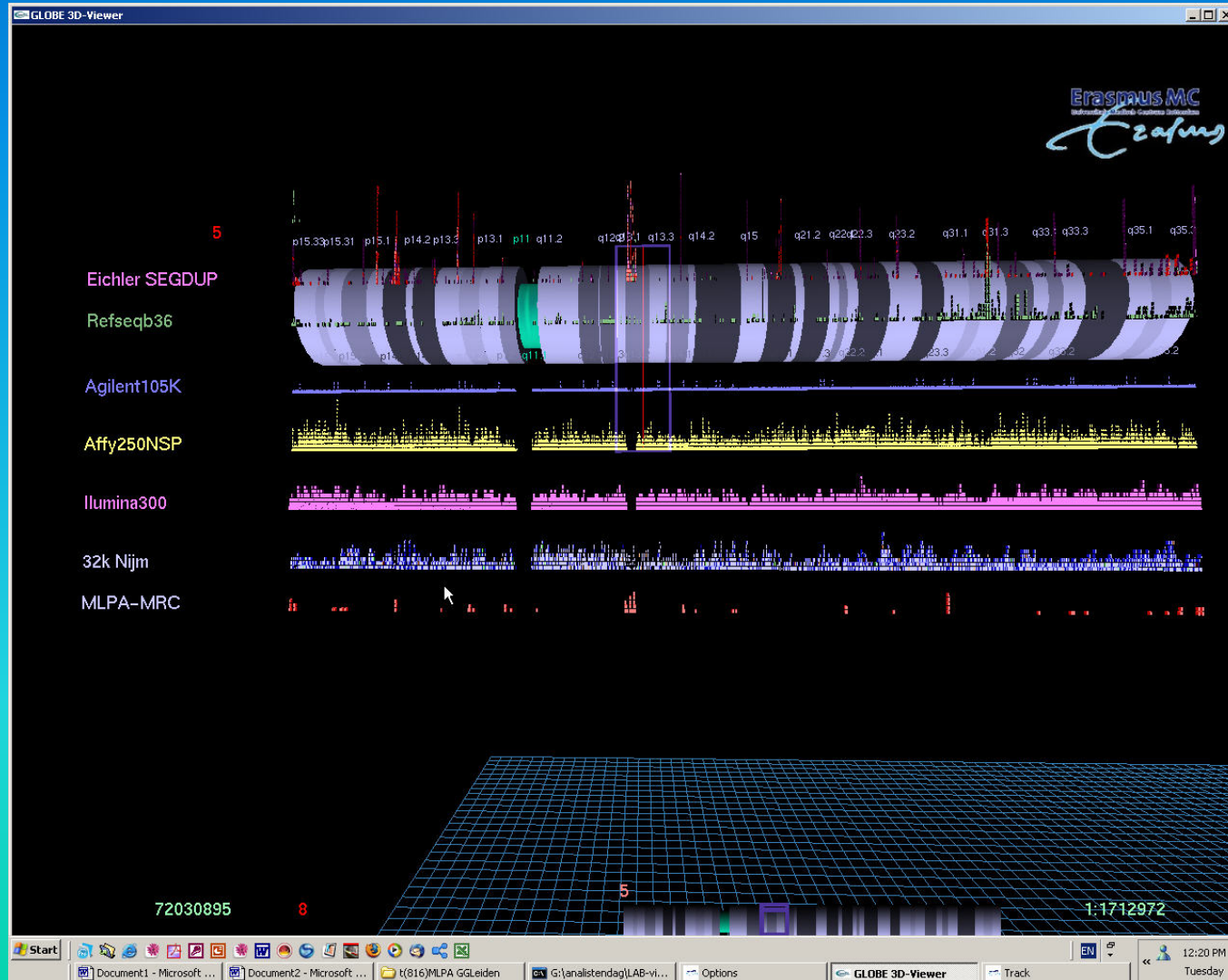
# Relations to Look at, are the Most Sexy Ones

The Human Visual System is Still Better than any Program !



# The Neighbors Wife is Always Attractive!

Comparison of Different Data/Experiments is the Key in Genomics !



# The Devils Sitting on the Cake

## Current Solutions in the Genomic Sector

### Properties of Current Solutions

- difficult to use and long training
- non generic and home-brewn
- integration from base pair to chromosome

### The Witches

- content organizations: Ensemble, UCSD
- home made solutions and small business
- non-classical competitors: Google, Adobe

### Our Wisdom Team

- > 20 years of experience in genomics
- 3 expertiese covering 100% of genomics
- knowledge at the scientific and clinical frontier

### >>> We outperform by:

- knowledge leadership
- technological leadership

The screenshot displays the Ensembl Human CytoView interface. At the top, it shows the search bar and navigation options. The main content area is titled 'Chromosome 15' and shows a detailed view of a genomic region. The interface includes a 'Detailed view' section with various tracks for genomic features, including genes, transcripts, and repeats. A legend at the bottom identifies the different feature types, such as 'Ensembl Known Protein Coding', 'Vega External Known Protein Coding', and 'Repeats'. The interface also includes an 'Export data' section with options for selecting features, format, and type to export.



# The Cave of the Gold and Diamonds

## The Market Opportunities in the Genomic Sector

“I want this tool: NOW”

Evin Eichler  
Big Shot in Genomics  
Washington

### General Market Size

- > 100,000,000 species
- > 7,000,000,000 individuals
- > 1,000,000 disease relevant

“Every CEO wants to have it on a USB”

ENZO  
general product manager

### Research and Development Sector

- ~500,000 genomic workers
- >10,000 academic institutes
- >10,000 BioTech / Pharma R&D

### Commercial Sector

- ~500,000 diagnostic applicators
- >100,000 diagnostic Labs
- >5,000 BioTech/Pharma Suppliers

“You are 10 years ahead of the field”

Viviane Reding  
EU Commissioner  
Information Society & Media

- Working in genomics centres
- Pharma R&D
- Euros per year

### In The Netherlands

- >2000 diagnostic workers
- >100 diagnostic labs
- >300 diagnostic companies present
- >100,000 clinical cases per year

>>> The Market has Huge Growth Potential > 50 years !  
>>> The Genomic Sector will not loose its importance !



# The Gold and The Diamonds

## Financial Opportunities and Return of Investments



### *Market Opportunities*

- licenses
- white label products
- content sale
- advertising
- e-commerce

### *Individual R&D licence*

- academic: 1,000 Euro/a
- industry: 5,000 - 20,000 Euro/a

### *Corporate R&D licence*

- academic: 50,000 - 200,000 Euro/a
- industry: 100,000 - 200,000 Euro/a

### *Genomic Product Suppliers*

- sales portal: 100,000 - 200,000 Euro
- sales share: 2 - 5 % of turnover

### *Diagnostic Institutions*

- individual licence: 3,000-7,000 Euro/a

### *We have already:*

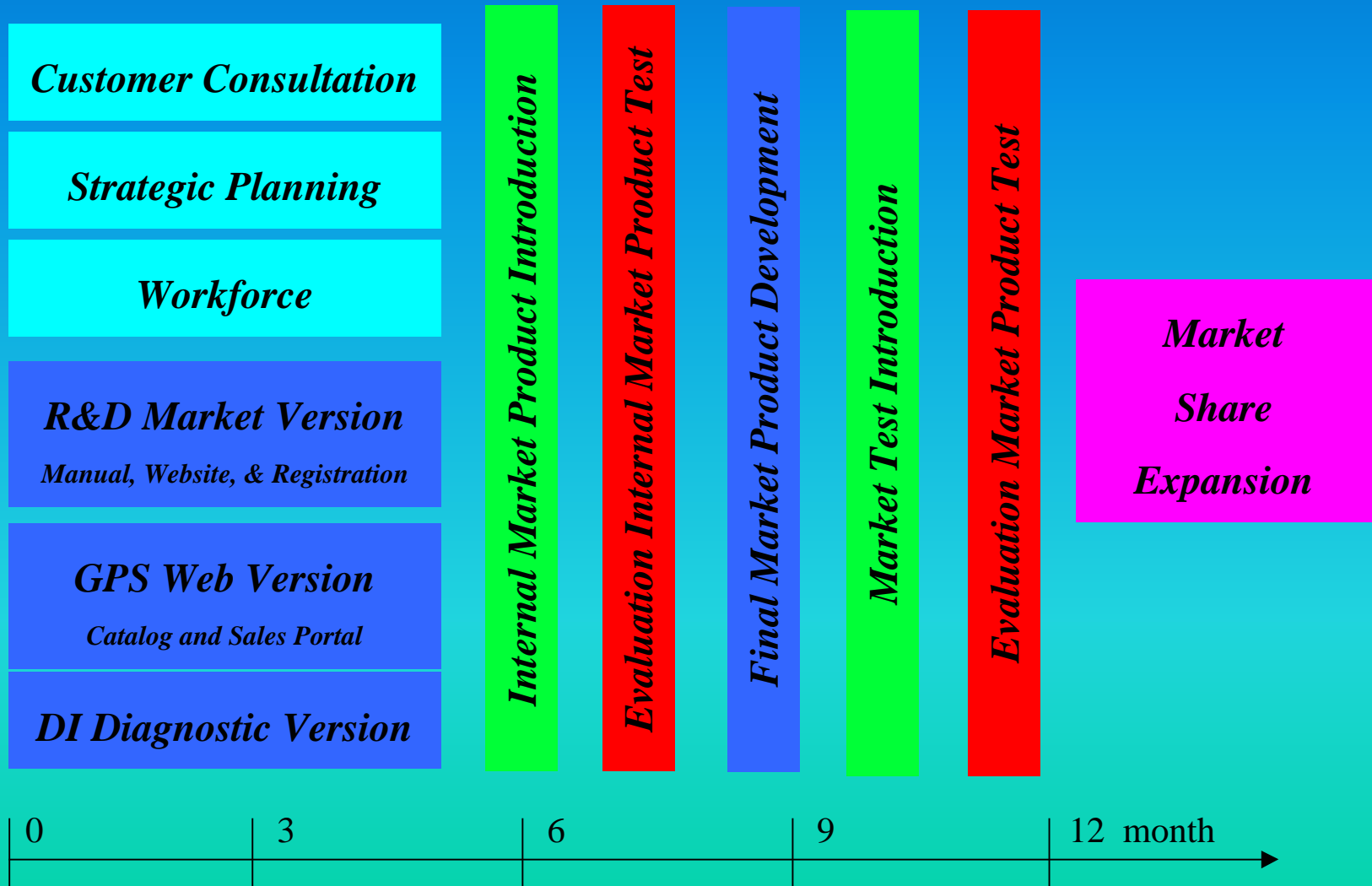
- >10 Erasmus MC internal customers (on a cooperation basis)
- >10 cooperation users world wide (on a cooperation basis)
- the German MediGRID as a corporation partner
- Erasmus MC Incubator management financing: 20,000 + 80,000 Euro



# It is a **SHORT** Way to Tipperary

## 12 Month Plan for a (Test) Market Introduction

Budget: For the **WHOLE** Process Cheap 200 kEuro.



# The GLOBE 3D Genome Browser

**Knoch, T. A., Eussen, B. & Moorhouse, M.**

*NBIC Venture Challenge, University of Neijmegen, Neijmegen, The Netherlands,  
19th June, 2007.*

## *Abstract*

The combination of genome sequence and structure, its annotation and experimental data in an accessible and comprehensible way is a major challenge. Increasingly, there are a large number of extremely divergent data sets: the sequence itself, genes, regulatory regions, various forms of reoccurring sequence features and clone sets etc. Currently, one possibility to represent this information in a visual form - and thus to reveal its scientific meaning - is to use genome browsers such as "Ensembl" or "The UCSC Genome Browser". These browsers have been beneficial in the understanding of the complex organization of genomes. However, there are also huge limitations concerning their focus on linear presentation, standardized input and data bank accessibility. Also customisability by a remote user with special requirements is difficult.

Therefore, the GLOBE-Consortium has developed the next generation genome viewer - the GLOBE 3D-Viewer - to visualize multi-dimensional data sets from various sources in an easy intuitive manor while accounting for the complexity of genome organization. Together with an easily accessible data-warehouse for archiving (un-)published experimental data and the Erasmus Computing Grid (ECG) for fast interactive large-scale (correlative) analysis the GLOBE-Consortium creates an intuitive holistic environment for genome research. Thus, the GLOBE 3D-Viewer sets the new standard to integrate complex genomic data sets within a single holistic display system and thus is major advance for scientific and clinical genome research.

### **Rational behind the GLOBE 3D-Viewer**

The GLOBE 3D-Viewer is based on many a basic rational driven by the complex nature of genome organization in general and by the diverse individual research and diagnostic data types in particular. Thus, the key features of the GLOBE 3D-Viewer are:

- \* flexibility, customisability, intuitive navigation
- \* real-time interaction and analysis
- \* dynamic scaling, semantic zooming, and object arrangement
- \* visualization of extremely large data sets and multi-dimensional data
- \* bridging all scales of genome organization from sequence to morphology

### **Capabilities of the Prototype GLOBE 3D-Viewer**

The GLOBE 3D-Viewer allows the integration of all possible linear and structural genomic types of information from the DNA sequence to complete genomes in a three-dimensional environment. Genomic elements are visualized in a flexible and customisable manor and can be arranged in relation to each other as desired. The navigation within this three-dimensional space is intuitive and resembles the virtual environments of flight simulators or architectural planning tools. This supports also rapid resolution change, i.e. zooming-out from the base pair to the whole-genome level is possible within an eye-glimpse using fly-in/out paradigm while including

appearance or relational positioning changes of genetic elements. The GLOBE 3D-Viewer is capable of relating extremely large numbers of genomic elements and multi-dimensional data-sets to each other with inbuilt analysis running in real-time.

In addition, the GLOBE 3D-Viewer is able to present the three-dimensional organisation of genomes from base pair resolution to complete chromosomes. This allows the direct projection of linear genomic information on its real three-dimensional spatial architecture for the first time.

The GLOBE 3D-Viewer integrates e.g. the following genomic and experimental elements or data types in a three-dimensional virtual environment:

- \* Genomic information types: Syndromes, break-points, duplicons, repeat regions, epigenetic annotations, genes and single nucleotide polymorphisms.
- \* Structural genome features: Chromosomes, ideogram bands, chromatin loops, chromatin fiber conformation, nucleosomes, DNA double helix.
- \* Experimental genome features: Bacterial Artificial Chromosomes, Yeast Artificial Chromosomes, Fosmids, genomic and proteomic arrays, restriction sites, oligomers, primers.
- \* Genomic experiment types: 3D fluorescence *in-situ* hybridization (3D-FISH), Metaphase fluorescence *in-situ* hybridization (M-FISH), comparative genome hybridisation (CGH), expression profiling, quantitative polymerase chain reaction (QPCR).

The GLOBE 3D-Viewer creates entire new possibilities in the visualization in the holistic properties of genomes. It opens new perspectives for future research leading to a better understanding of genomes and their system biological aspects. This is of major importance in respect to advanced diagnostics and ultimate disease treatments.

### **Opportunities of the GLOBE 3D-Viewer**

The GLOBE 3D-Viewer creates a new intuitive virtual environment to show for the first the enormous complexity of genomes in a *single display*. This leads to an all encompassing understanding of genomes and opens exciting new opportunities in every aspect of scientific, clinical, education as well as commercial applications:

- \* Scientists benefit from easier planning and experimental analysis as well as hypothesis building and better comprehension of genomic function.
- \* Diagnostic users benefit from a better overview of their diagnostic tasks and the evaluation of the results including patient counselling and treatment.
- \* Educators and publishers benefit from enhanced presentation of genome organisation and function in classrooms or e-publications.
- \* Commercial applicants benefit from improved planning of new products, their in test, increased sales as well as more appealing marketing of their products.

In summary, the GLOBE 3D-Viewer offers the opportunity to understand genomes on the necessary system-biological level for the first time, and thus sets the new standard for genome browsers for scientific and clinical research and as well as for unseen possibilities concerning educational and commercial purposes. Therefore, the GLOBE 3D-Viewer creates also unrivalled chances for world-wide introduction generating a sustainable revenue stream.

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### Keywords:

Genome, genomics, genome organization, genome architecture, structural sequencing, architectural sequencing, systems genomics, coevolution, holistic genetics, genome mechanics, genome function, genetics, gene



regulation, replication, transcription, repair, homologous recombination, simultaneous co-transfection, cell division, mitosis, metaphase, interphase, cell nucleus, nuclear structure, nuclear organization, chromatin density distribution, nuclear morphology, chromosome territories, subchromosomal domains, chromatin loop aggregates, chromatin rosettes, chromatin loops, chromatin fibre, chromatin density, persistence length, spatial distance measurement, histones, H1.0, H2A, H2B, H3, H4, mH2A1.2, DNA sequence, complete sequenced genomes, molecular transport, obstructed diffusion, anomalous diffusion, percolation, long-range correlations, fractal analysis, scaling analysis, exact yard-stick dimension, box-counting dimension, lacunarity dimension, local nuclear dimension, nuclear diffuseness, parallel super computing, grid computing, volunteer computing, Brownian Dynamics, Monte Carlo, fluorescence in situ hybridization, confocal laser scanning microscopy, fluorescence correlation spectroscopy, super resolution microscopy, spatial precision distance microscopy, auto-fluorescent proteins, CFP, GFP, YFP, DsRed, fusion protein, in vivo labelling, information browser, visual data base access, holistic viewing system, integrative data management, extreme visualization, three-dimensional virtual environment, virtual paper tool.

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