

51. Jahrestagung der Deutschen Gesellschaft für
Medizinische Informatik, Biometrie und Epidemiologie (gmds)

10. - 14. September 2006

Spatial Genetic Heterogeneity Assessment - a regional case study

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Genetic Heterogeneity

Genetic epidemiology

Evolution

Clinical studies

Pharmacogenetics

allele frequency

F-statistics

cluster analysis

wombling

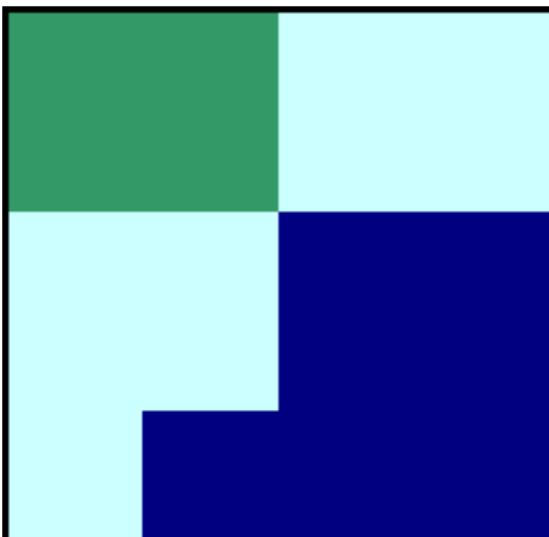
principal component analysis

spatial structure

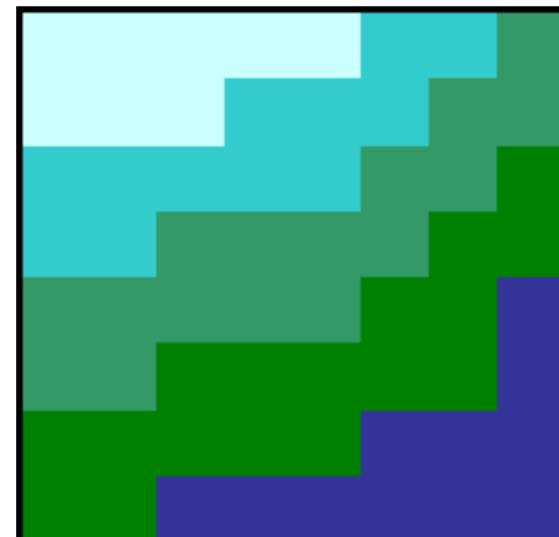
Genetic landscapes

Genetic landscapes

Clusters



Clines



Genetic Landscapes

KORA S4

Probands

- 700 probands
- ethnicity and socioeconomic information

Markers

- randomly selected markers
- 212 SNPs
- autosomal and X-chromosomal
- 1:2 intragene / intergene position

Steffens et. al., 2006, SNP-Bases Analysis of Genetic Substructure in the German Population, *Hum Hered* (in press)

http://www.gsf.de/KORA/projekt_survey2000.html



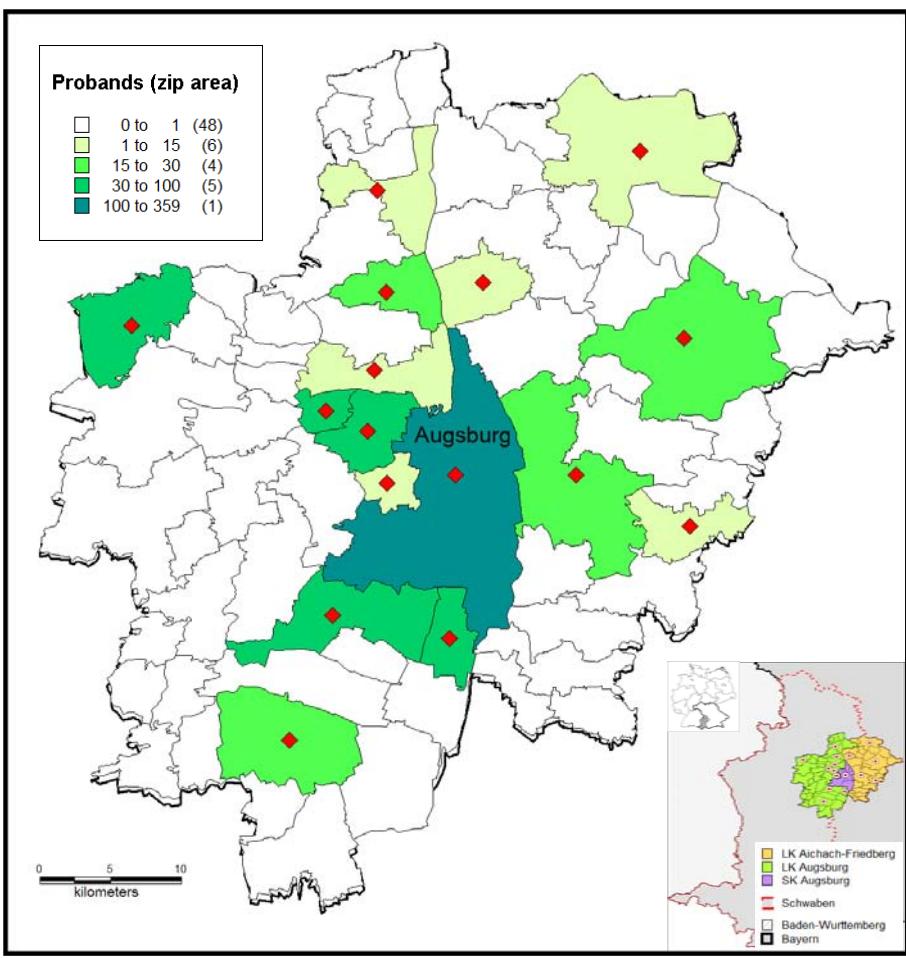
GRASS

Geographic Resources Analysis Support System

Spatial modelling and analysis

of genetic information

KORA S4



Data

726 Probands
29 zip areas
187 SNPs

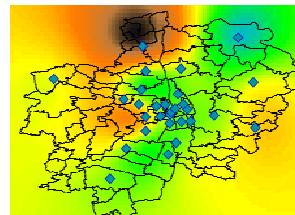
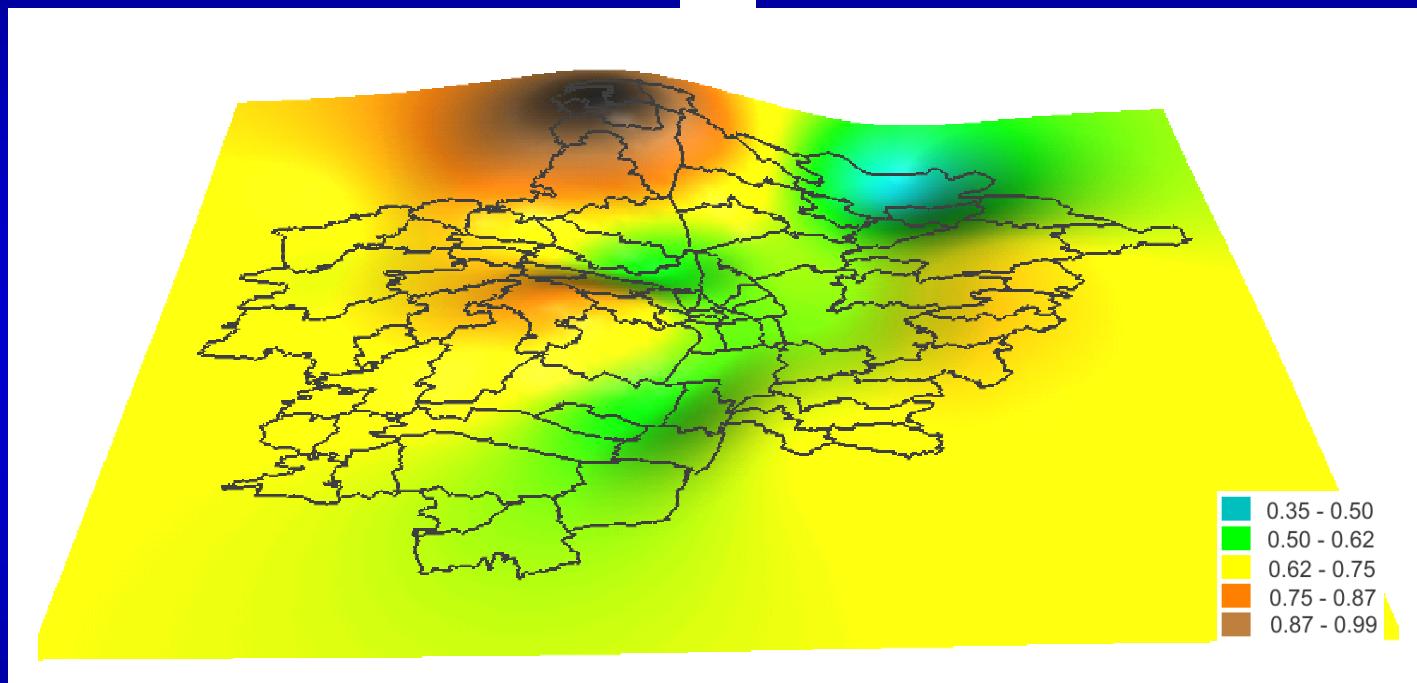
- count (minor allele)
- $q = \text{minor allele frequency}$
-> $f = \arcsin(\sqrt(q))$

Spatial Differentiation of Allele Frequency

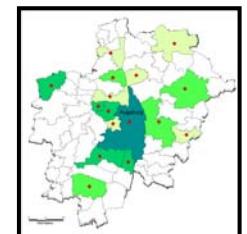
| rs3625 (count) | | |
|----------------|---------|---------|
| PLZ | ALLELE1 | ALLELE2 |
| 86150 | 324 | 364 |
| 86316 | 20 | 24 |
| 86343 | 41 | 41 |
| 86356 | 36 | 52 |
| 86368 | 8 | 2 |
| 86391 | 5 | 7 |
| 86399 | 48 | 54 |
| 86405 | 8 | 14 |
| 86450 | 51 | 45 |
| 86462 | 14 | 30 |
| 86482 | 33 | 29 |
| 86495 | 6 | 10 |
| 86508 | 11 | 11 |
| 86551 | 31 | 11 |
| 86554 | 9 | 13 |
| 86830 | 15 | 17 |

| MARKER | p-value | Postion (gene) |
|-----------------|---------------|----------------|
| rs1860300 | 0,0062 | inter |
| rs1997660 | 0,0324 | intra |
| rs2242046 | 0,0174 | intra |
| rs3625 | 0,0138 | intra |
| rs4379869 | 0,0631 | intra |
| rs461311 | 0,0128 | inter |
| rs597354 | 0,0076 | inter |
| rs717477 | 0,0018 | inter |

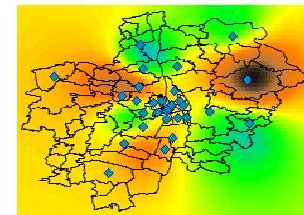
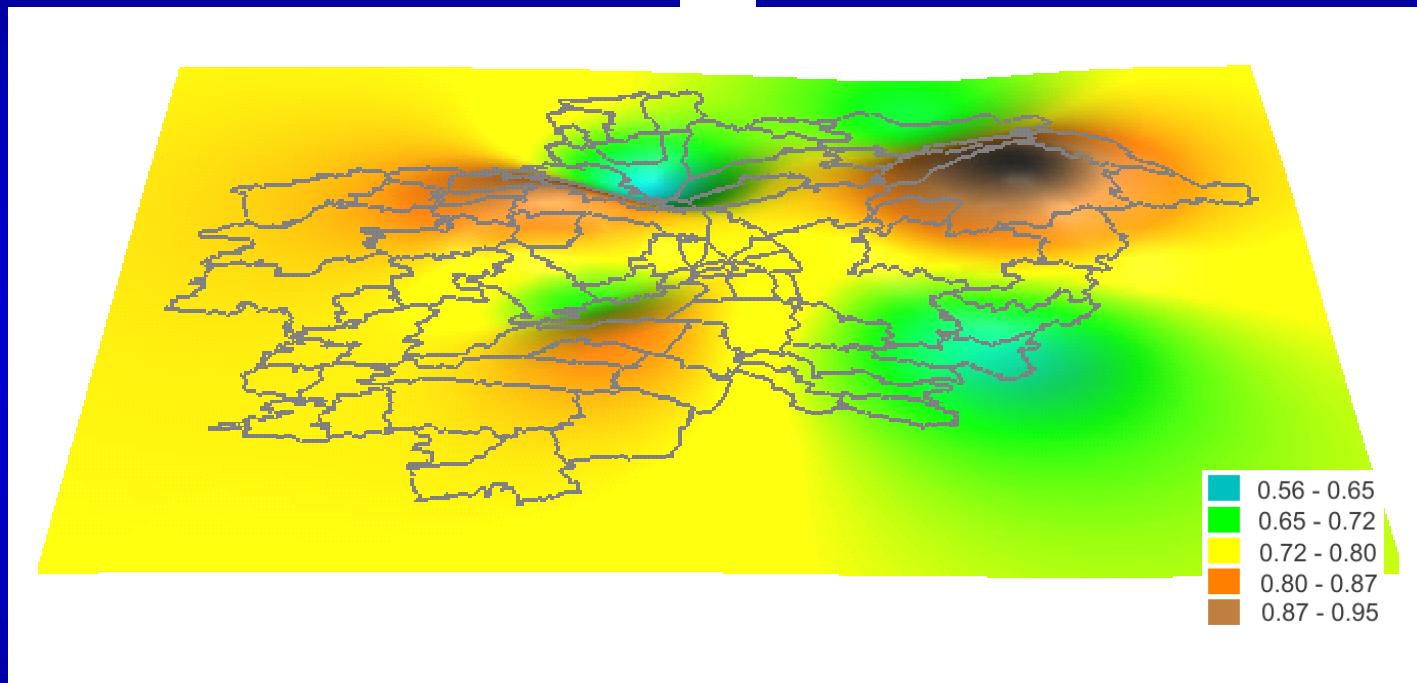
Genetic Landscapes



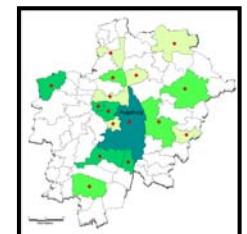
Allele Frequencies - rs717477



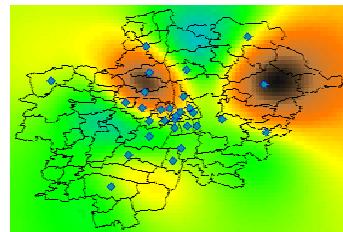
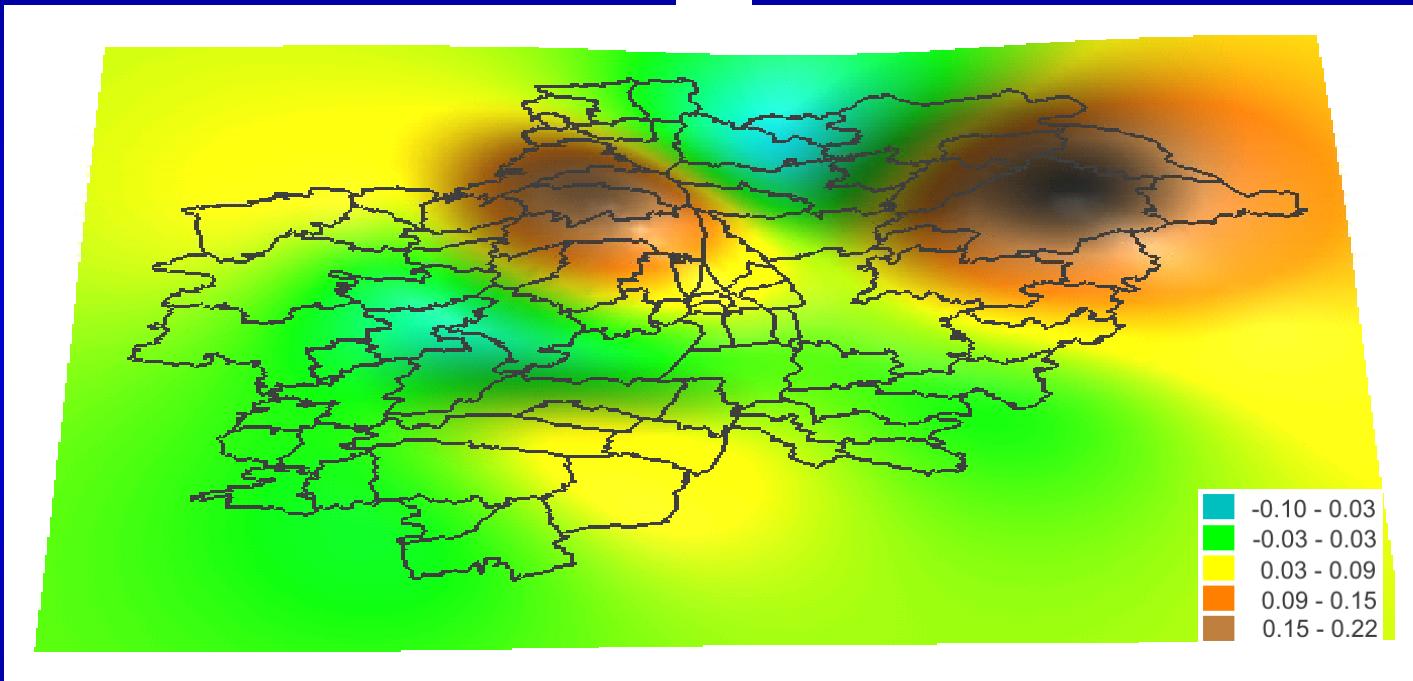
Genetic Landscapes



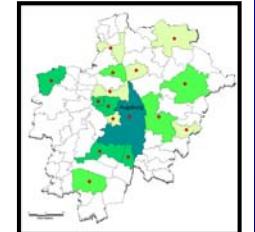
Allele Frequencies - rs3625



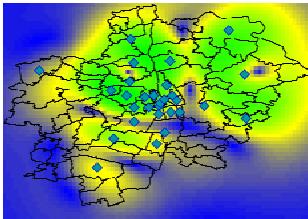
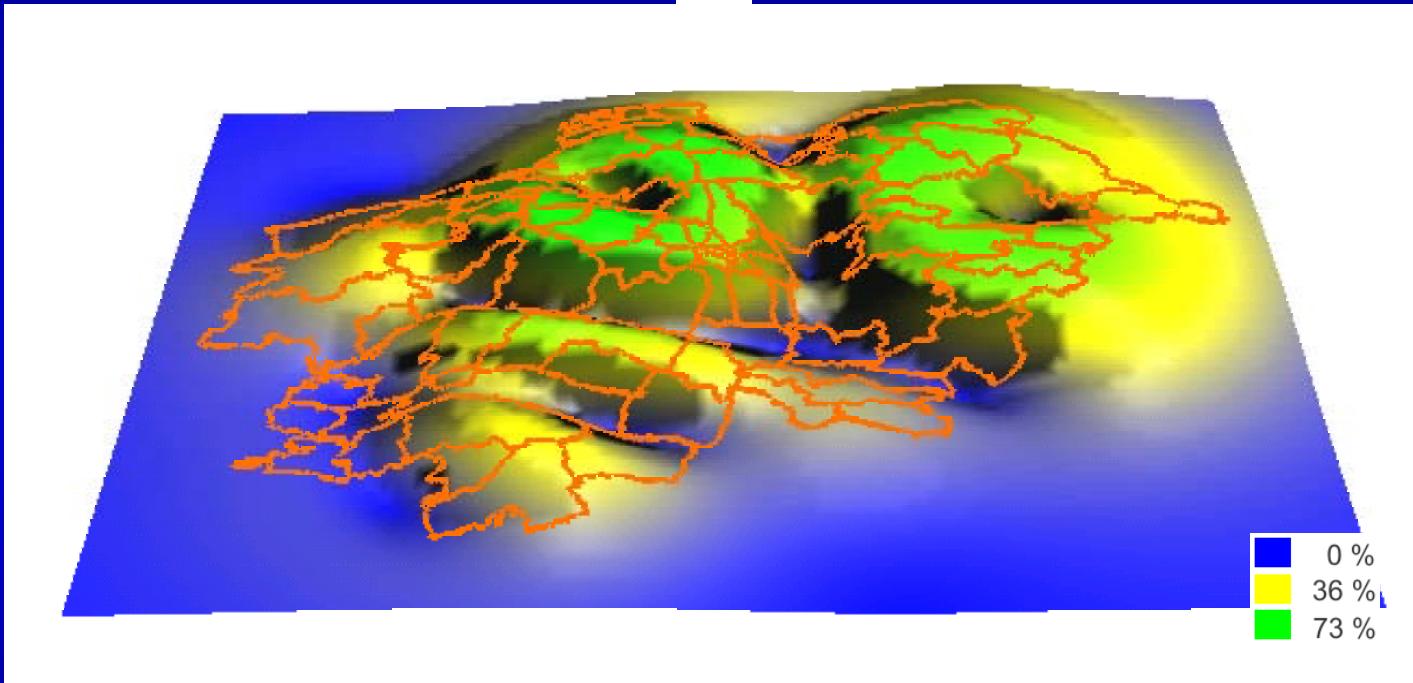
Genetic Landscapes



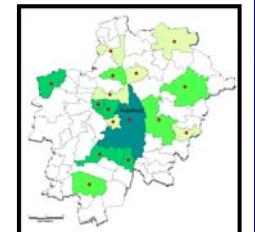
Population Specific F_{IS} - rs3625



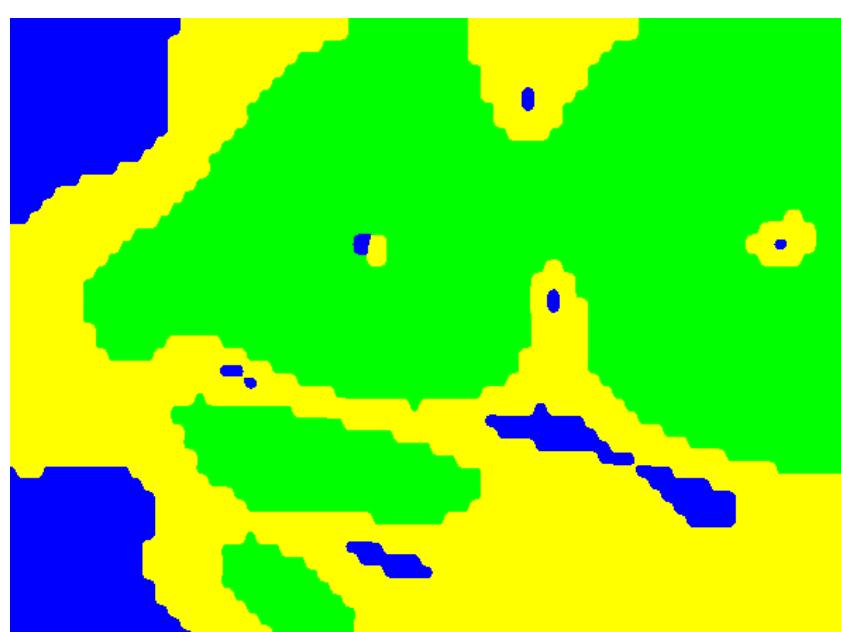
Genetic Landscapes



Slope of Population spc. F_{IS} - rs3625

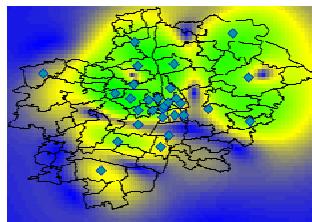


Genetic Landscapes

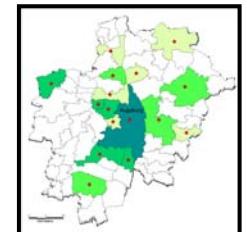


| Class | Slope (%) | Pixel |
|-------|----------------|-------|
| 1 | Low < 10 | 100 |
| 2 | Medium 10 - 30 | 633 |
| 3 | High ≥ 30 | 2501 |

| Landscape Index | Score |
|--------------------------|---------------|
| Patchiness (1:2:3) | high (10:3:3) |
| Connectivity (class: 1) | low |
| Fragmentation (class: 1) | high |



Slope of Population spc. F_{IS} - rs3625



Conclusions

A Genetic Geographical Information System is a powerful tool for detecting, modelling and analysing genetic heterogeneity along a study region.

Spatially limited populations may contain significant genetic heterogeneity for a single allele.

The widespread assumption of homogeneity by genetic studies should be evaluated case specifically.

Acknowledgments

ITC

Trento

Markus Neteler

Cesar Furlanello

gsf

Munich

Claudia Larnina

Erich Wichmann

IMBE

University of Bonn

Tim Becker

Michael Steffens

Arnfried Schiller

Thomas F. Wienker

*GenGIS opens the GATE to the
third dimension in genetic studies.*

...Thank you