ay/bi199: methods of computational science

visualization

visualization system+techniques

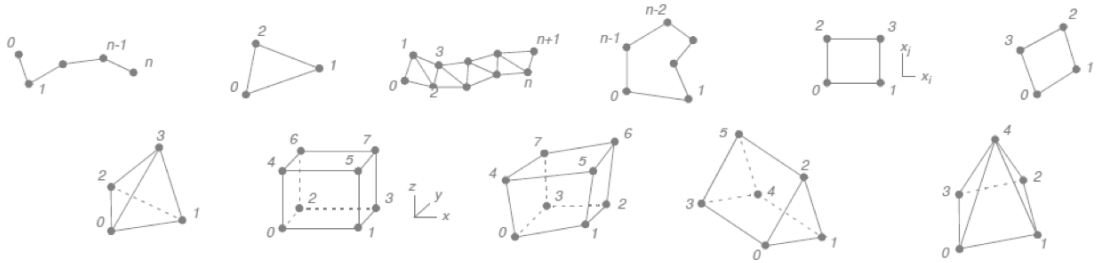
santiago v lombeyda | center for advanced computing research | caltech

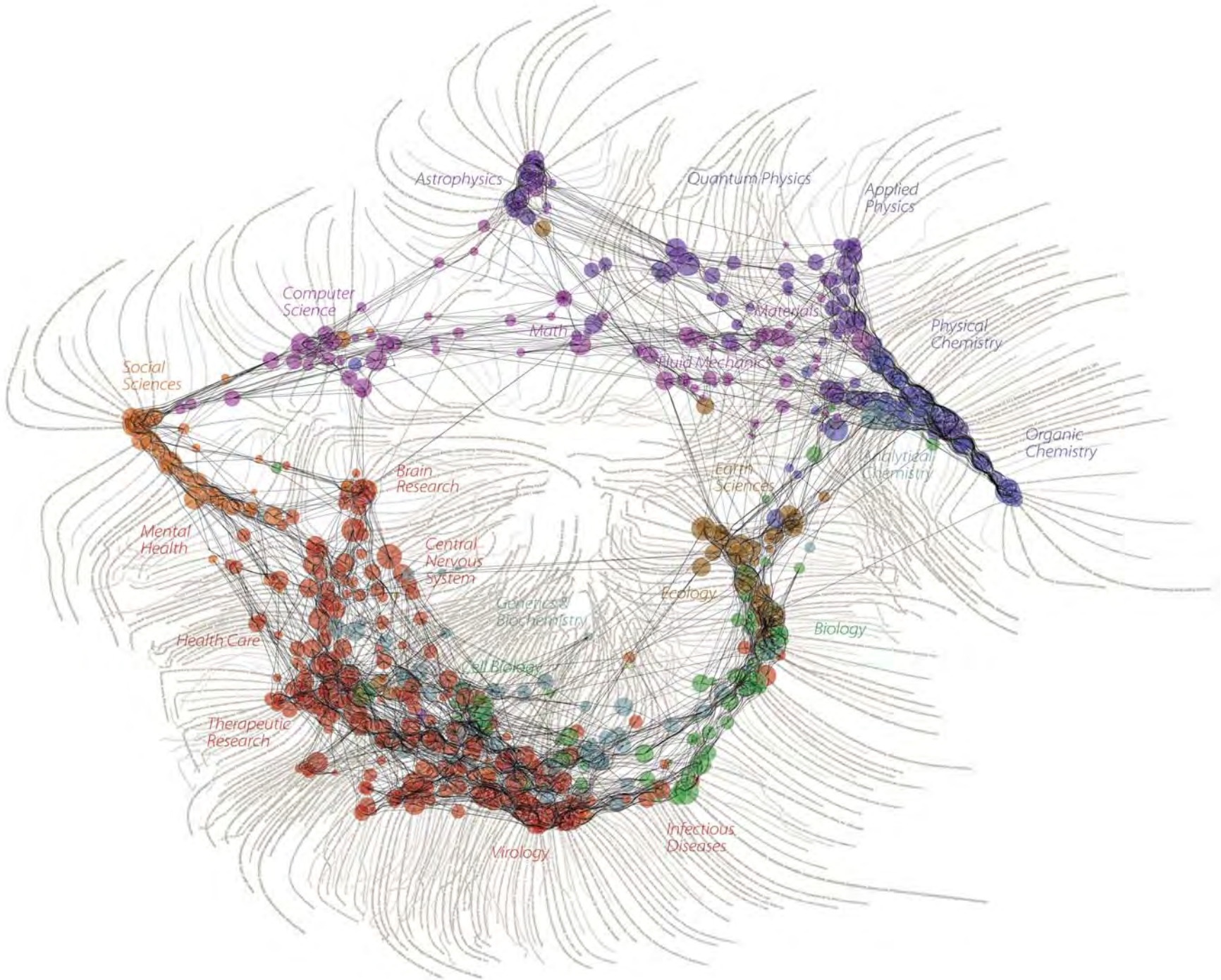
data: GEOMETRIC STRUCTURE

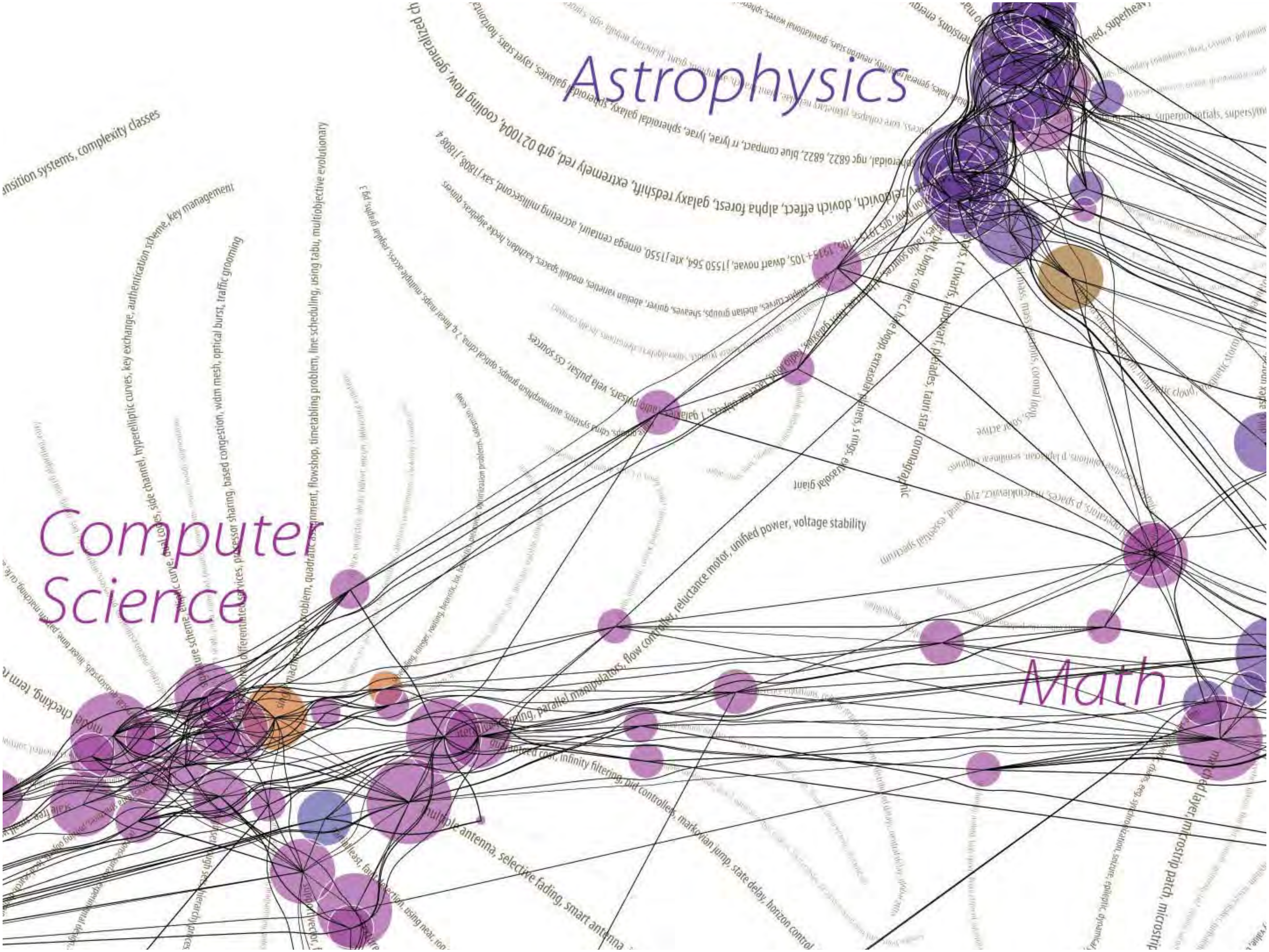
abstract

MPG Cylinders Horsepower Weight Acceleration Year Origin
8. 50.4 2.8 8.2 4 40. 250. 4 1500. 5500. 4 5. 30. 4 69.5 82.5 4 .8 3.2 3
18.000000 8.000000 130.000000 3504.000000 12.000000 70.000000 1.000000
15.000000 8.000000 165.000000 3693.000000 11.500000 70.000000 1.000000
18.000000 8.000000 150.000000 3436.000000 11.000000 70.000000 1.000000
16.000000 8.000000 150.000000 3433.000000 12.000000 70.000000 1.000000
17.000000 8.000000 140.000000 3449.000000 10.500000 70.000000 1.000000
... ...

2d/3d data




Astrophysics
Quantum Physics
Applied Physics
Computer Science
Materials
Physical Chemistry
Math
Fluid Mechanics
Social Sciences
Organic Chemistry
Analytical Chemistry
Brain Research
Earth Sciences
Mental Health
Central Nervous System
Ecology
Genetics & Biochemistry
Biology
Health Care
Cell Biology
Therapeutic Research
Infectious Diseases
Virology


Astrophysics
Computer Science
Math
unified power, voltage stability
transition systems, complexity classes
multiple antenna, selective fading, smart antenna
guaranteed cost, infinity filtering, pid controllers, markovian jump, state delay, horizon control


CHEMISTRY
MATERIALS
PSYCHOLOGY
ECONOMICS
MATHEMATICS
COMPUTER SCIENCE
GASTROENTEROLOGY
PSYCHIATRY
PARASITOLOGY
ECOLOGY & EVOLUTION
NEUROSCIENCE
MOLECULAR & CELL BIOLOGY
MEDICINE
GEOSCIENCES
MEDICAL IMAGING
ENVIRONMENTAL CHEMISTRY & MICROBIOLOGY
CROP SCIENCE
OTHER
FLUID MECHANICS
ANALYTIC CHEMISTRY
ASTRONOMY & ASTROPHYSICS
PHYSICS

& ASTROPHYSICS


ASTRON J
ASTRON ASTROPHYS
ASTROPHYS J
MON NOT R ASTRON SOC
ASTROPHYS J SUPPL S
ASTROPHYSICAL JOURNAL SUPPLEMENT SERIES
IN: 0.229497
OUT: 0.184635


ASTRON ASTROPHYS
ASTROPHYS J
MON NOT R ASTRON SOC
ASTROPHYS J SUPPL S

data: GEOMETRIC STRUCTURE

abstract

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16.000000 8.000000 150.000000 3433.000000 12.000000 70.000000 1.000000
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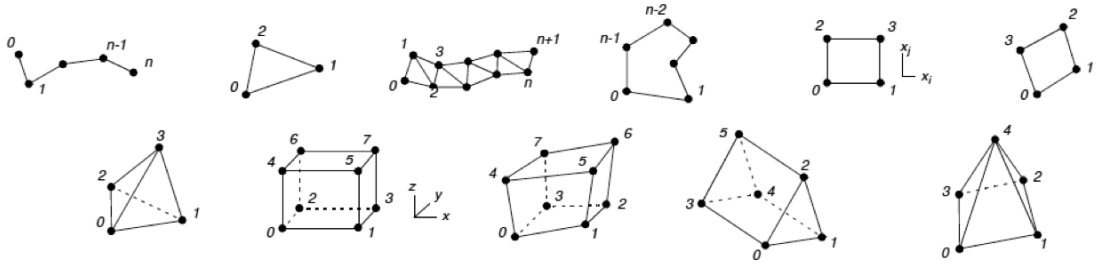
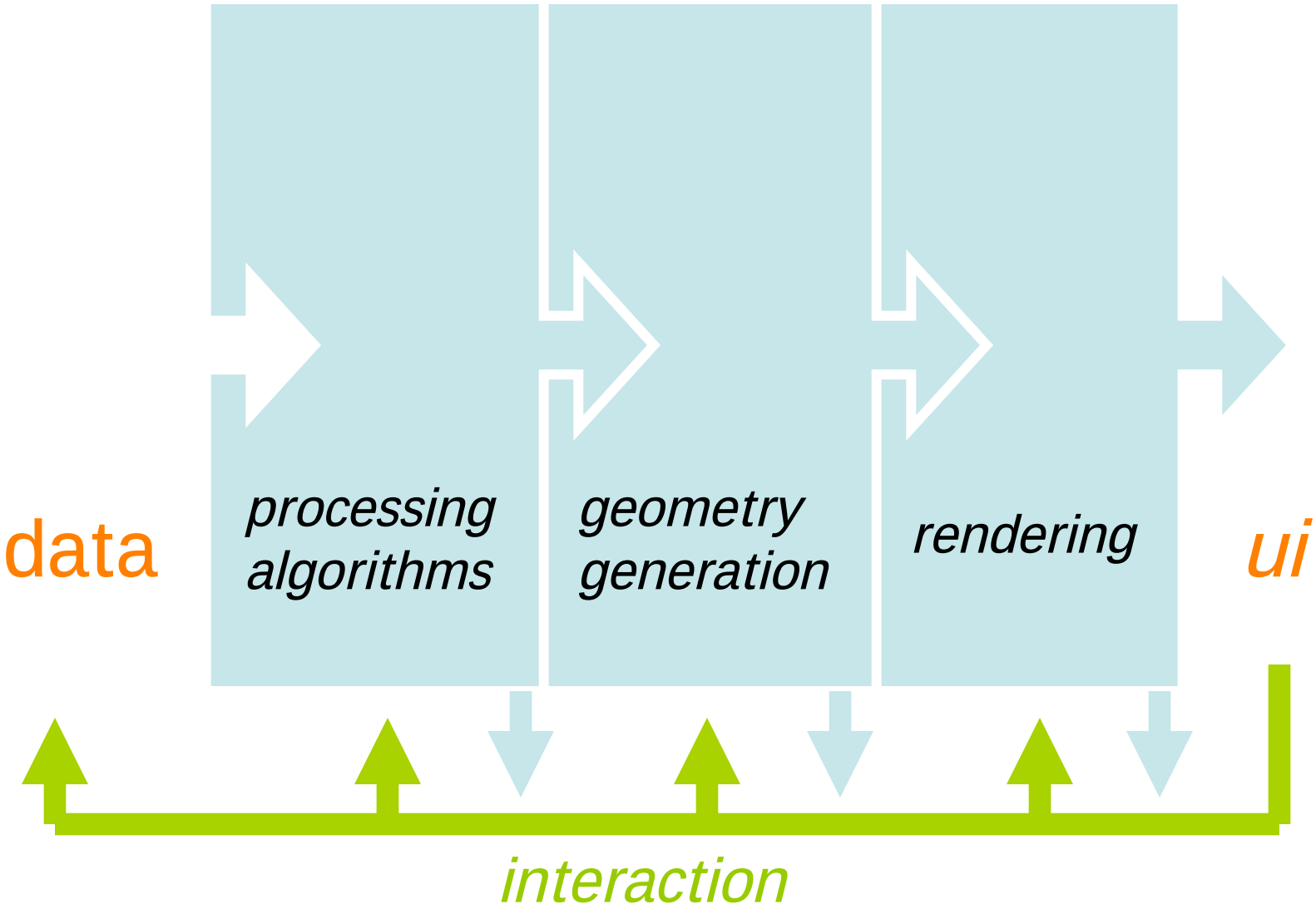
... ...

2d/3d data

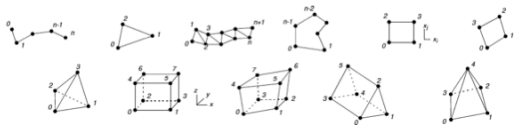


usual visualization engine

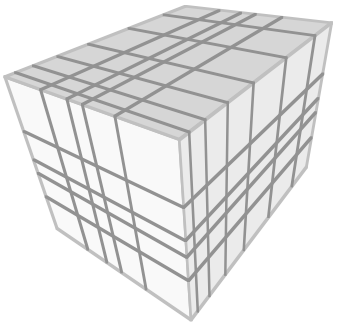


viz pipeline: *in the beginning*

DATA FORMATS



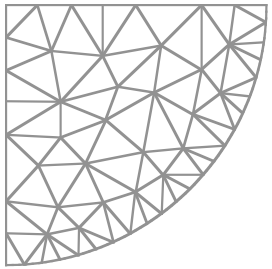
basic data types



structured grids

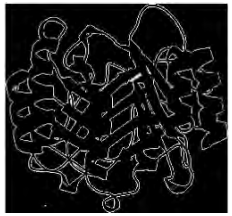
most basic: regular *(easy to do raw binary)*

more complex: AMR (adaptive mesh refinement)



unstructured grids

points
triangle meshes
tet meshes



atomic coordinate files

PDB (protein data bank)

no standard formats

raw

vtk

ascii or binary

new vtk

xml

ascii or base64 (mime) encoded binary

```
==   abcdefgh  ijklmnop
     qrstuvwx  yzABCDEF
     GHIJKLMN  OPQRSTUV
     WXYZ0123  456789+/
```

(64=2^6) 6 bits -> 8 bit char (*size * 4/3*)

amr

chombo(hdf5)/silo

pdb

standard..

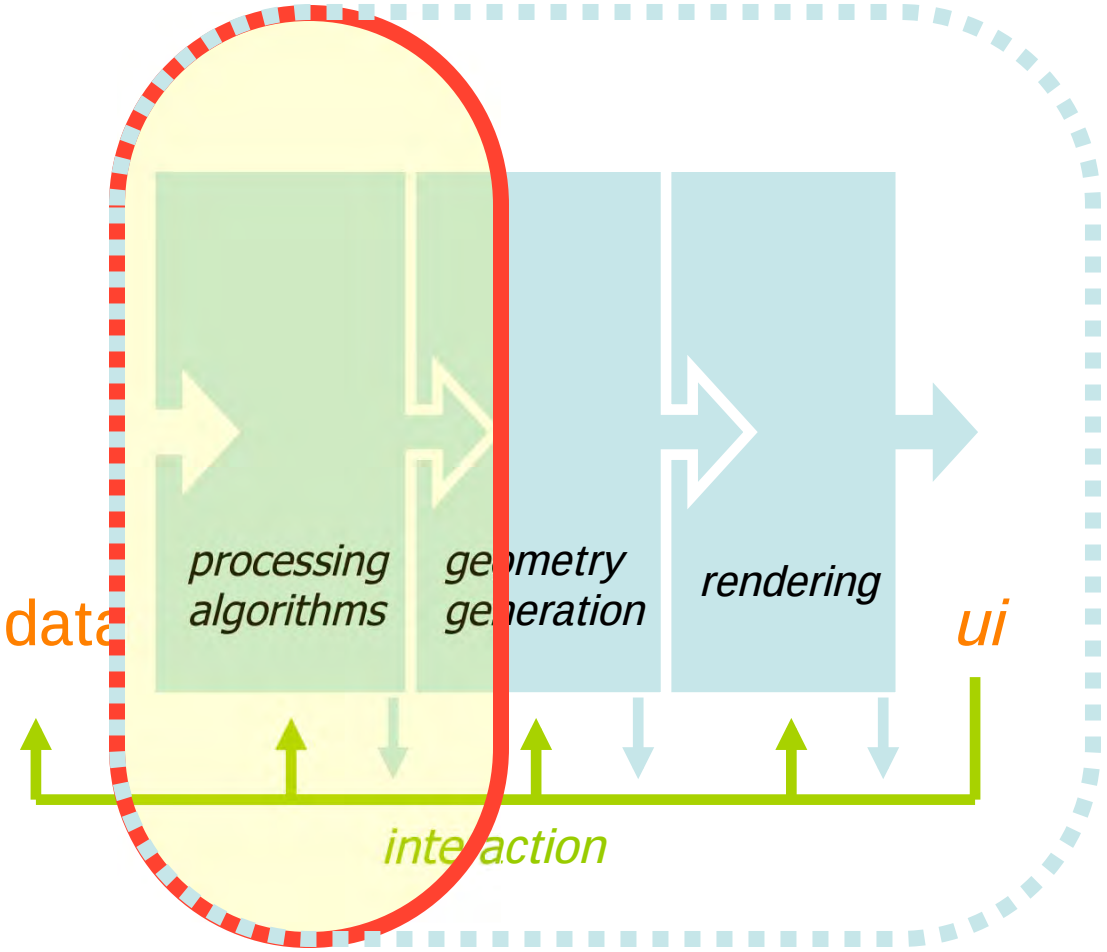
vtk sample: volume.vti

```
<?xml version="1.0"?>
<VTKFile type="ImageData" version="0.1" byte_order="LittleEndian">
<ImageData WholeExtent="0 3 0 3 0 3" Origin="0 0 0" Spacing="1 1 1">
 <Piece Extent="0 3 0 3 0 3">

  <PointData Scalars="vertexData">
   <DataArray type="Float32" Name="scalarData" format="ascii">
    0 1 2 3  1 2 3 4  2 3 4 8  3 6 9 11
    2 3 4 5  5 6 7 8  3 4 5 6  4 5 6 7
    3 4 5 6  3 4 5 6  4 5 6 7  6 7 8 9
    2 3 4 5  2 3 4 5  3 4 5 6  4 5 6 7
   </DataArray>
  </PointData>

  <CellData Scalars="cellData" Normals="cell_ normals">
   <DataArray type="Int32" Name="cellData" format="ascii">
    1 3 9  2 8 16  3 9 27
    2 3 4  6 7 8   6 9 10
    0 1 2  0 2 4   1 2 3
   </DataArray>
  </CellData>
 </Piece>
</ImageData>
</VTKFile>
```

“the” visualization toolkit



- VTK
  - c/c++
  - tcl/tk
  - python
  - java
- R

viz pipeline: *the tools*

VTK

the visualization toolkit

g.e. medical viz algorithms

➔ KITWARE

collection of filters

MARCHING CUBES *(patented)*

educational

➔ VTK book

evolved/extended

object oriented

C++

GL + Tk (UI)

now python

vtk

visualization algorithms

- scalar
- vector
- tensor
- texture
- volumetric

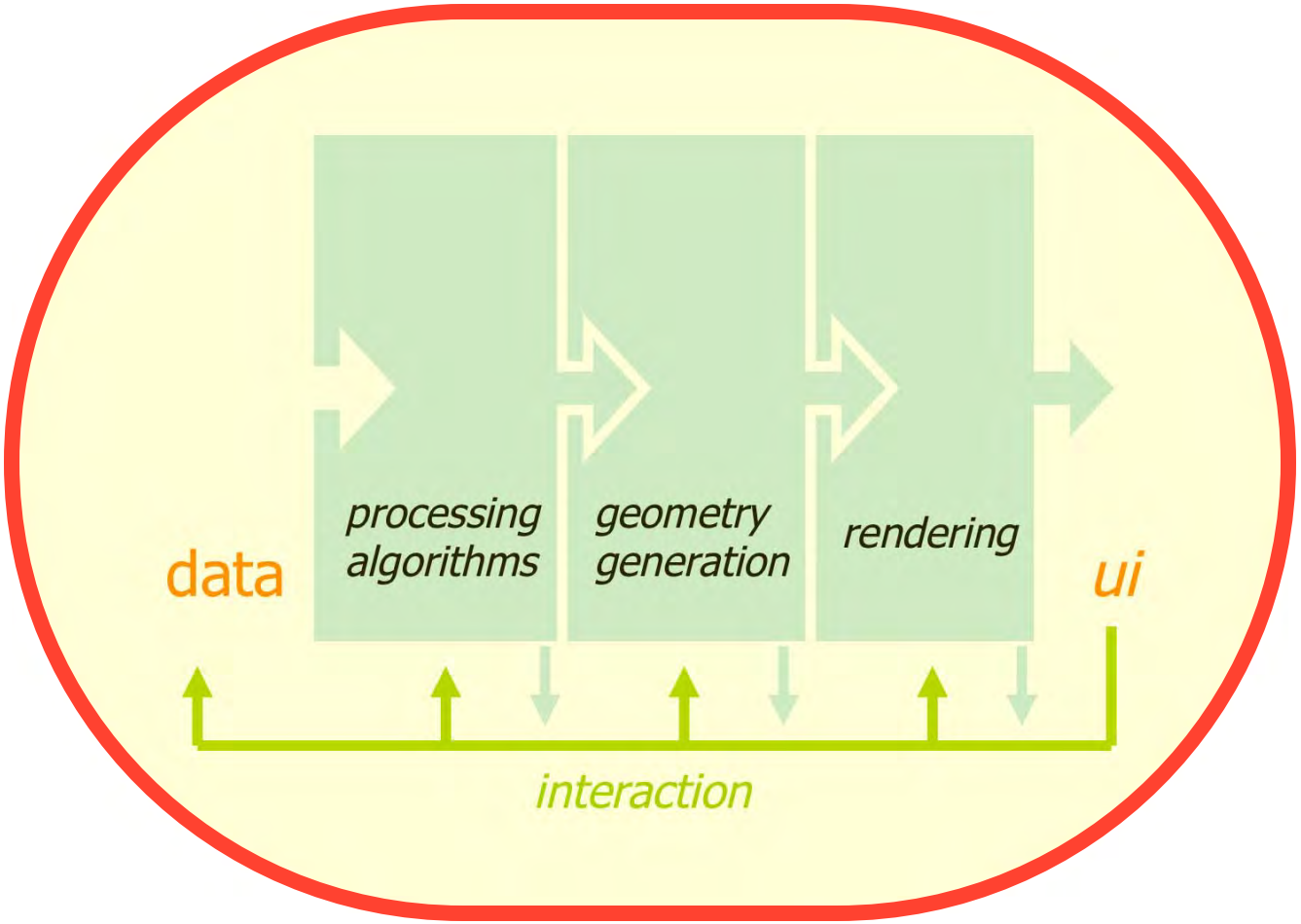
modeling techniques

- implicit modeling
- polygon reduction
- mesh smoothing
- cutting
- contouring
- Delaunay triangulation

imaging algorithms

- directly integrated
- mix 2D imaging/3D graphics

visualization system



- Paraview[VTK]
- LLNL VisIt[VTK]
- EnSight[$]
- Protovis[www]
- Many Eyes[www]
  - *Wiki based*
- Modrian[R],

  TomCat,

  Mollegro, ...

paraview

paraview.org

vtk based!

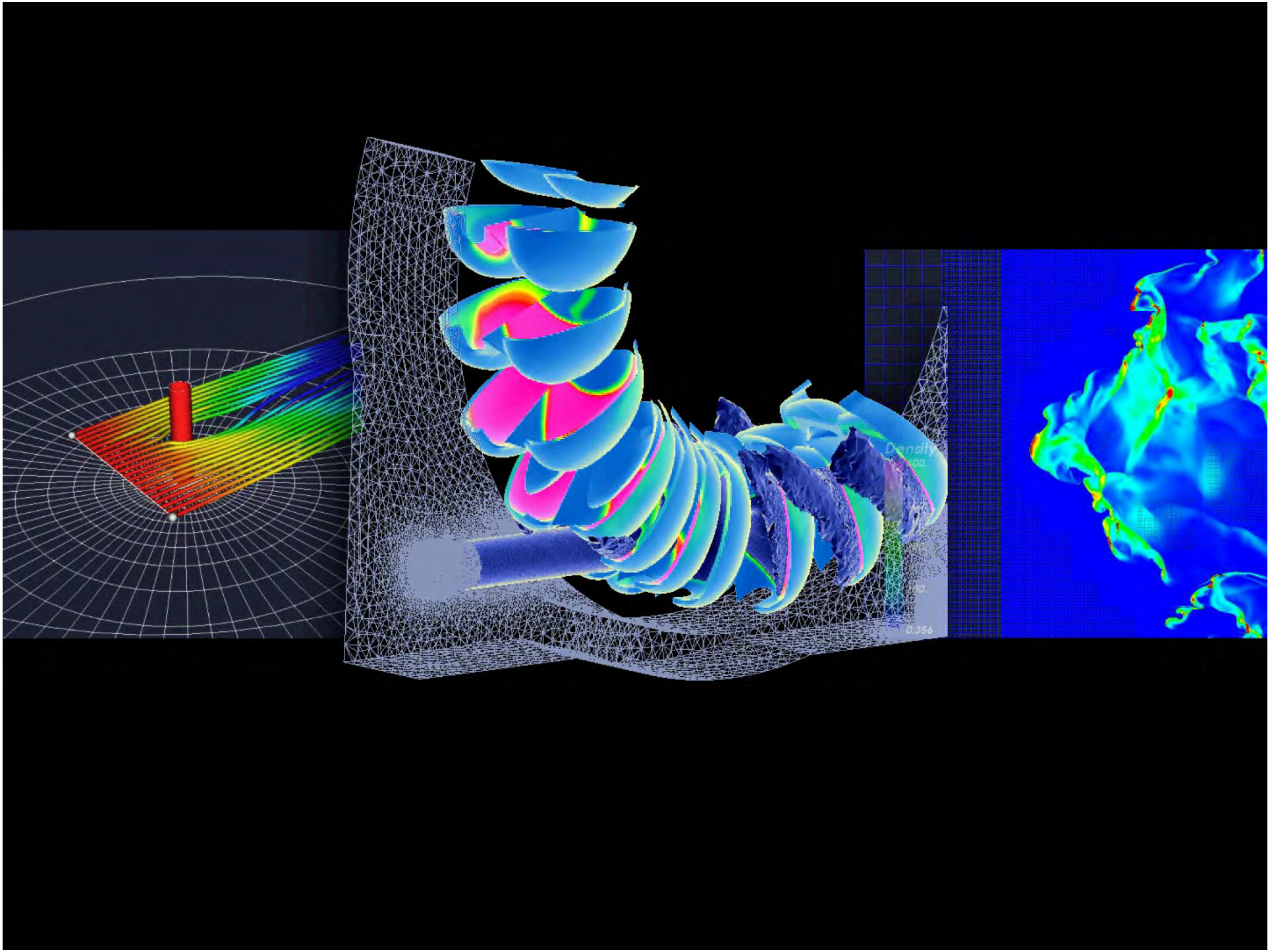
active community

- mailing list + wiki
- lead at kitware: *berk geveci*
  - sandia national lab
  - los alamos national lab, army research lab

parallel!

QT based

- starting 3.6.2 decent python scripting support

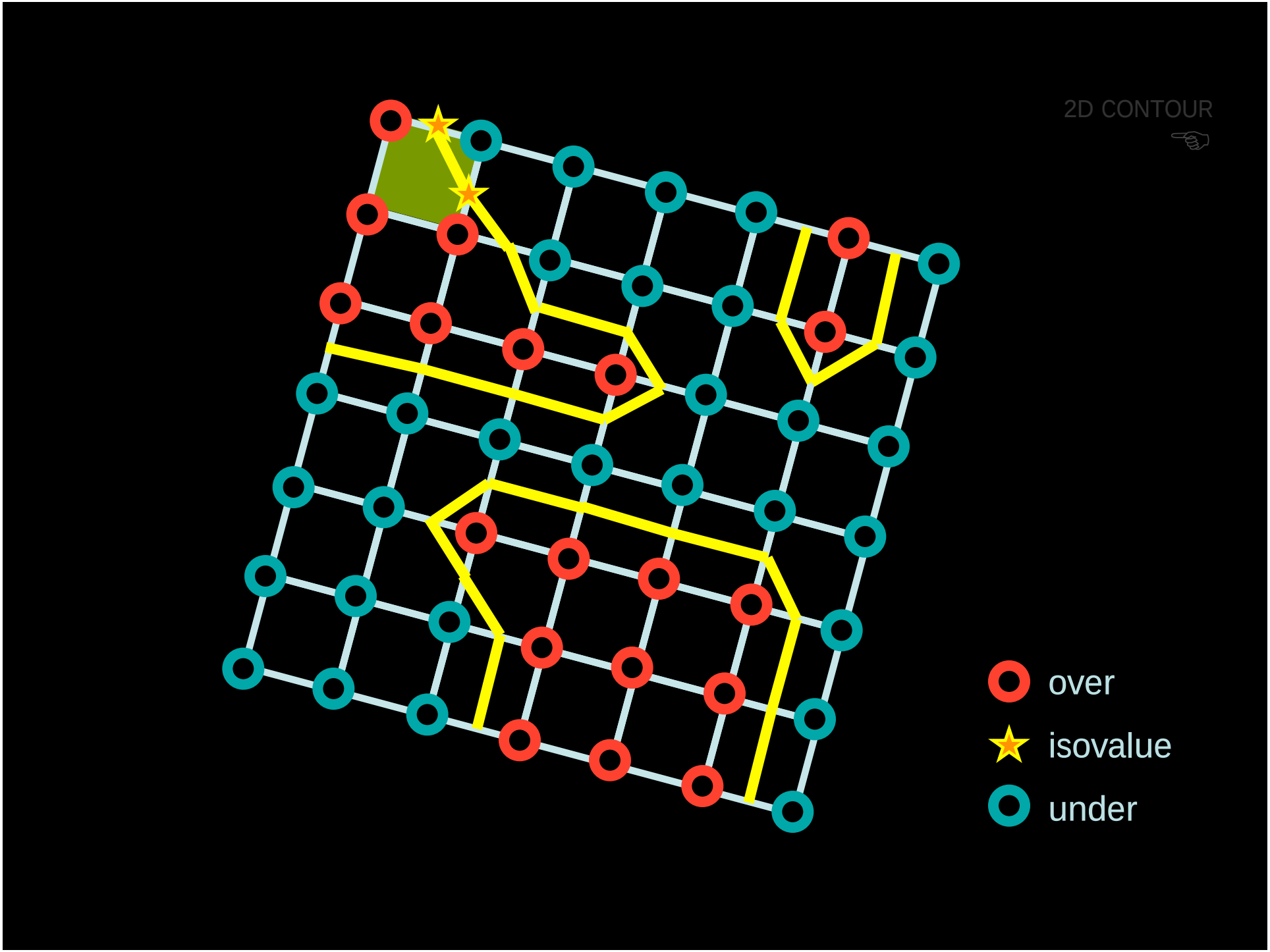

Density
0.356

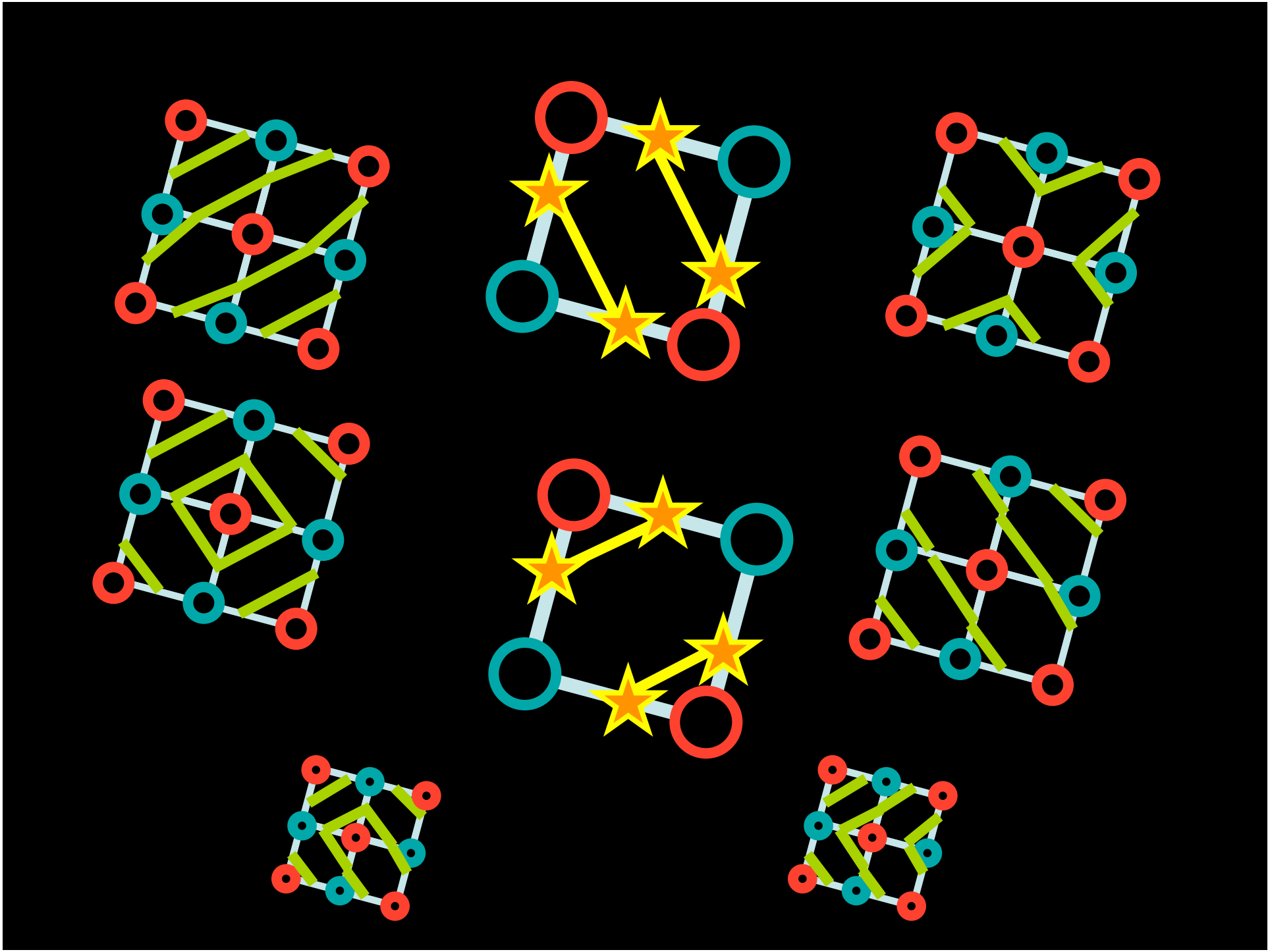
bluntfin:
PARAVIEW

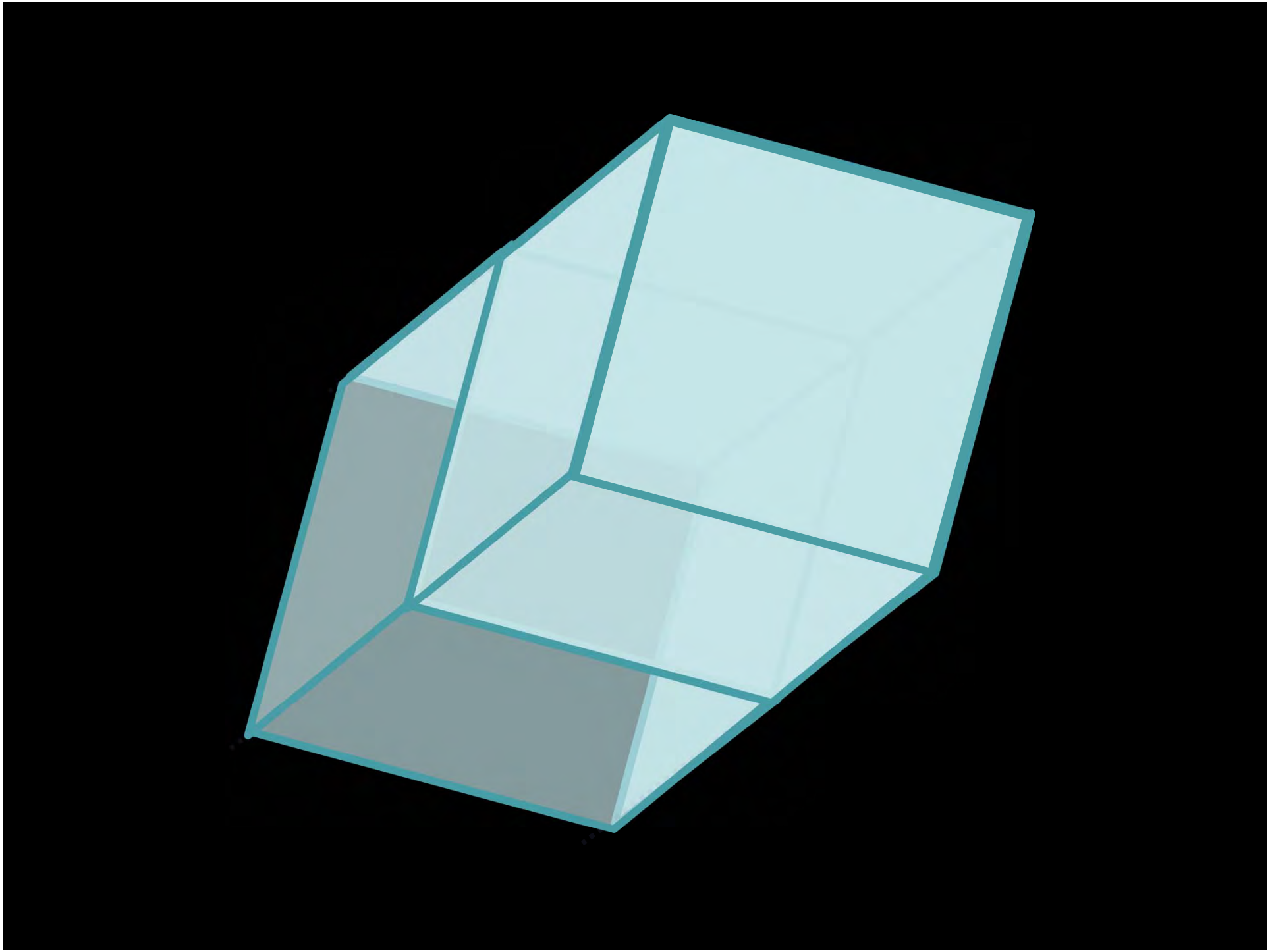
quick look

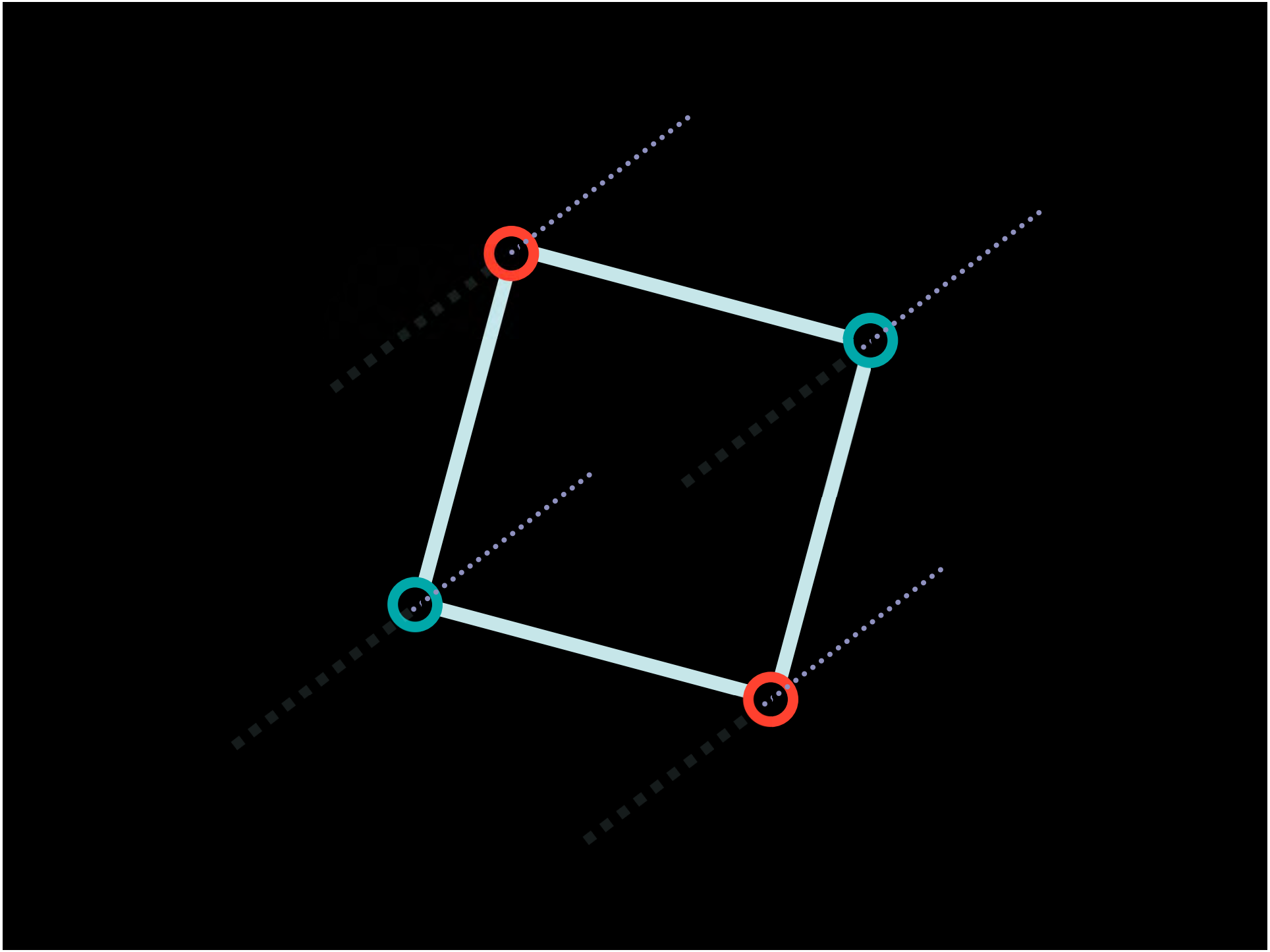
quick inside look: *marching cubes*

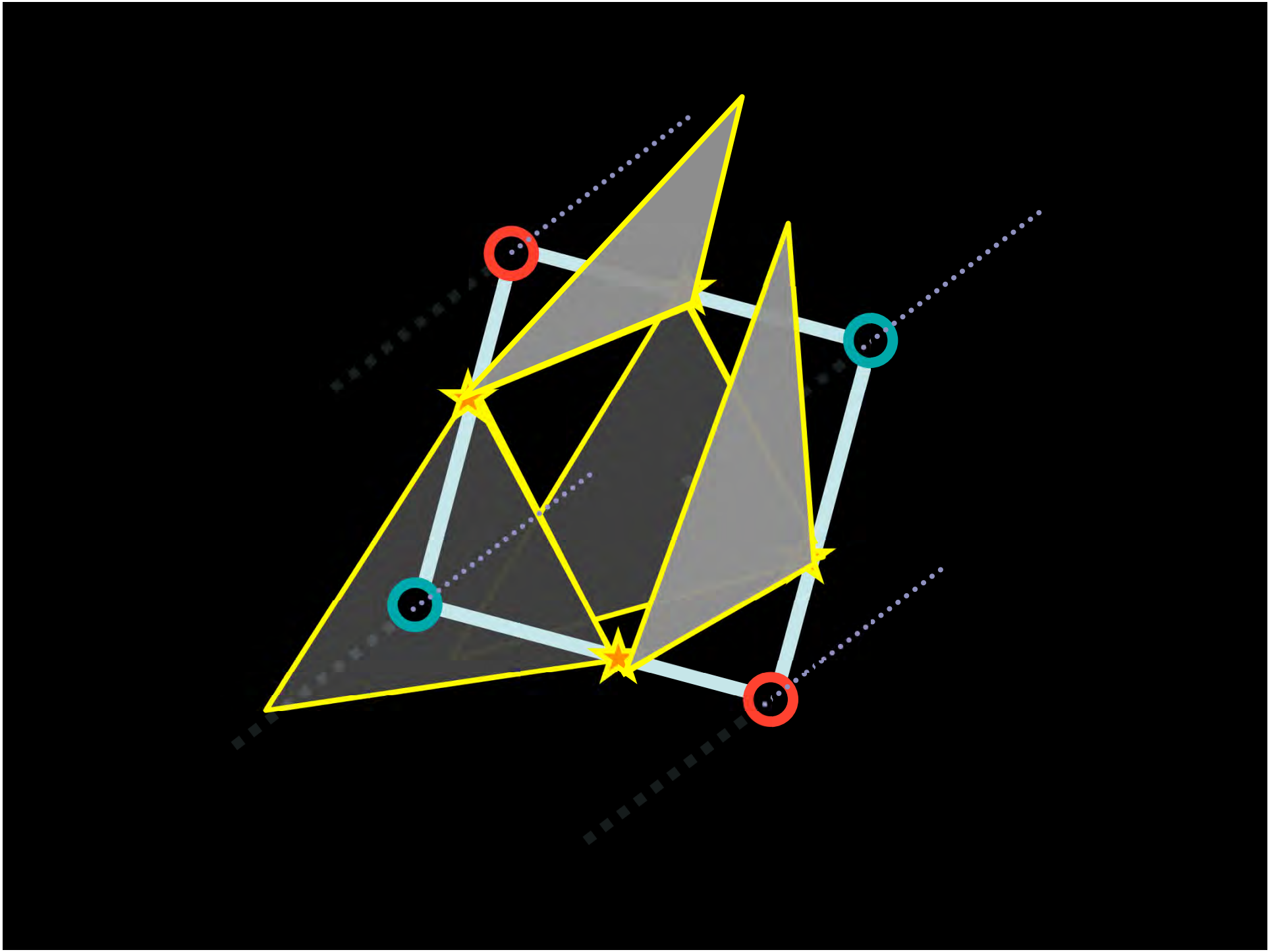
VTK

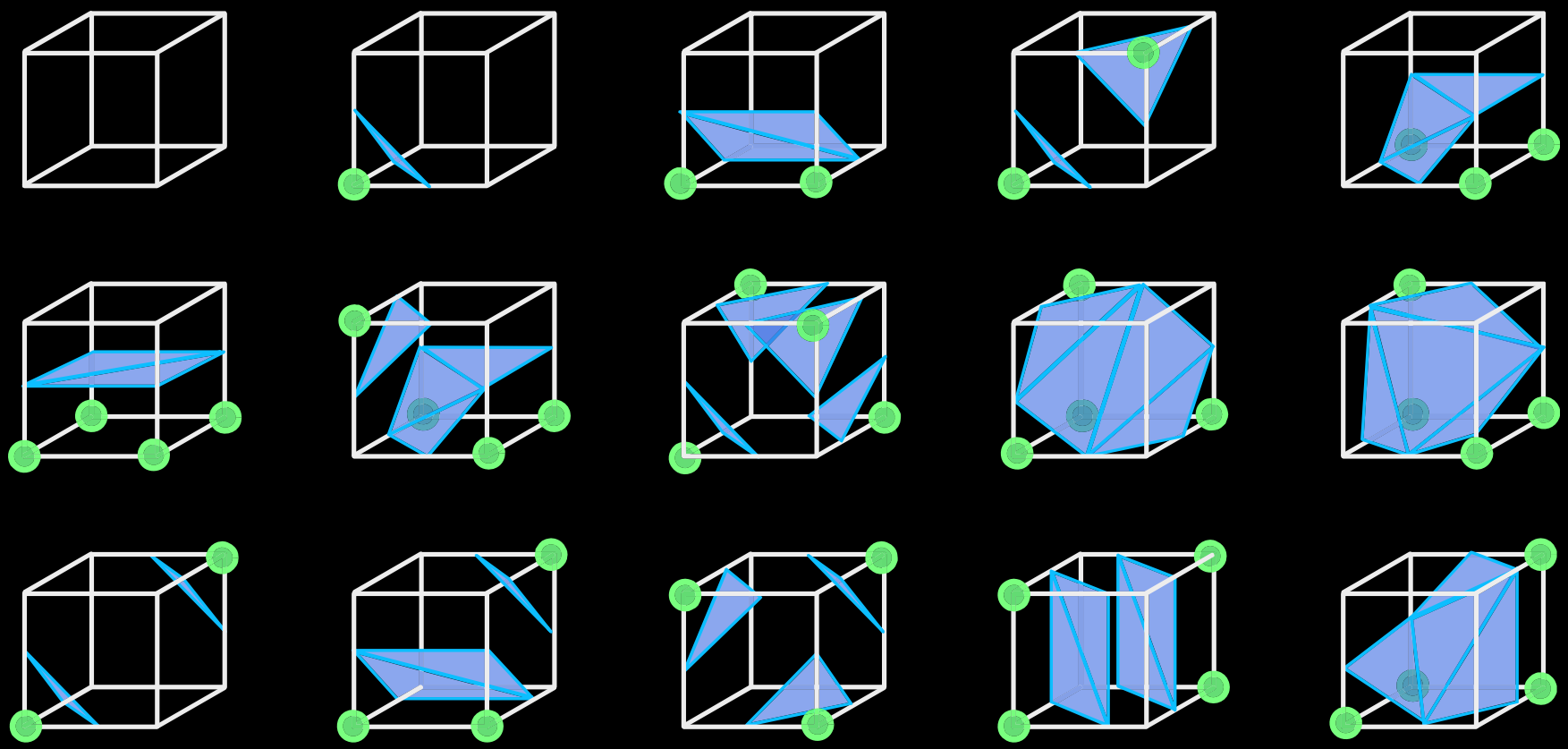

2D CONTOUR
over
isovalue
under



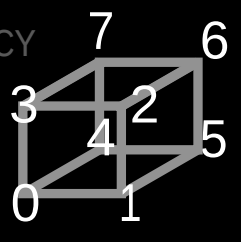








INCONSISTENCY ERROR FIX: TABLE FORCING CONSISTENCY



quick look: *techniques*

VOLUME RENDERING

project stargate:
NATIONAL LABS+SDSC

quick movie

volume rendering

VOLUMETRIC DATA

voxels: volume elements

usually regular 3D grid

scalar field

volume rendering

projected semi-transparent representation

scalar field + color transfer function (R,G,B,α)

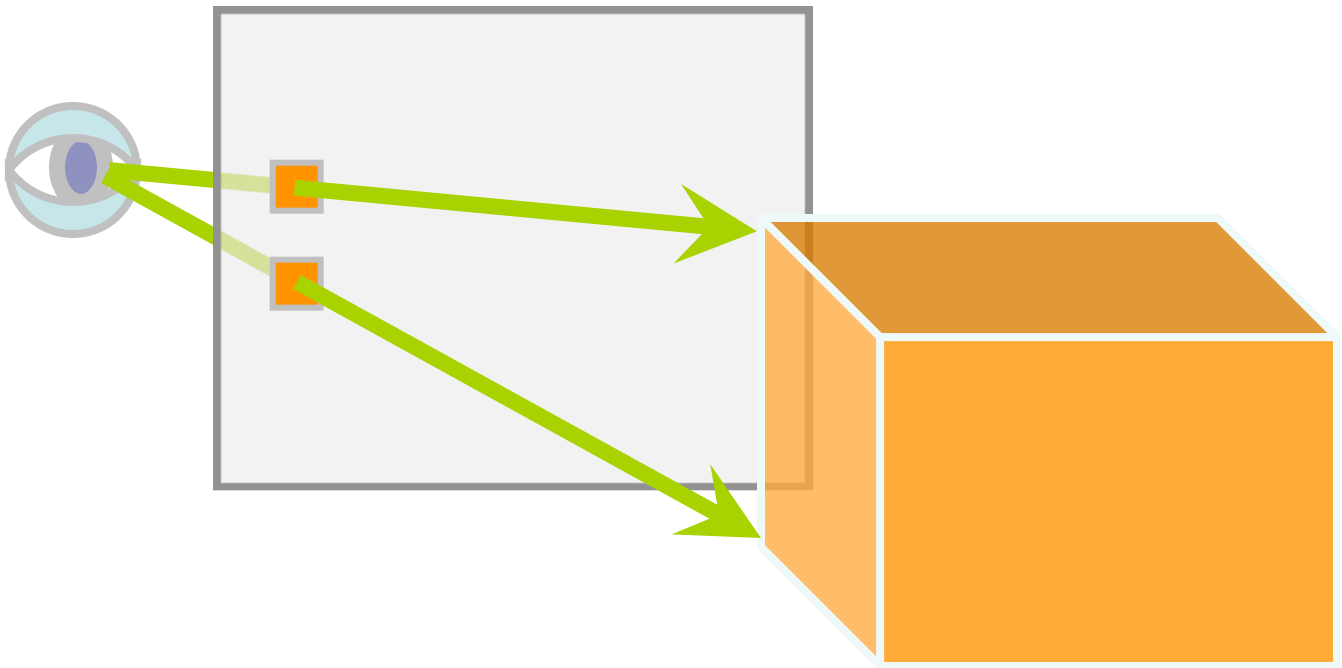
pixels

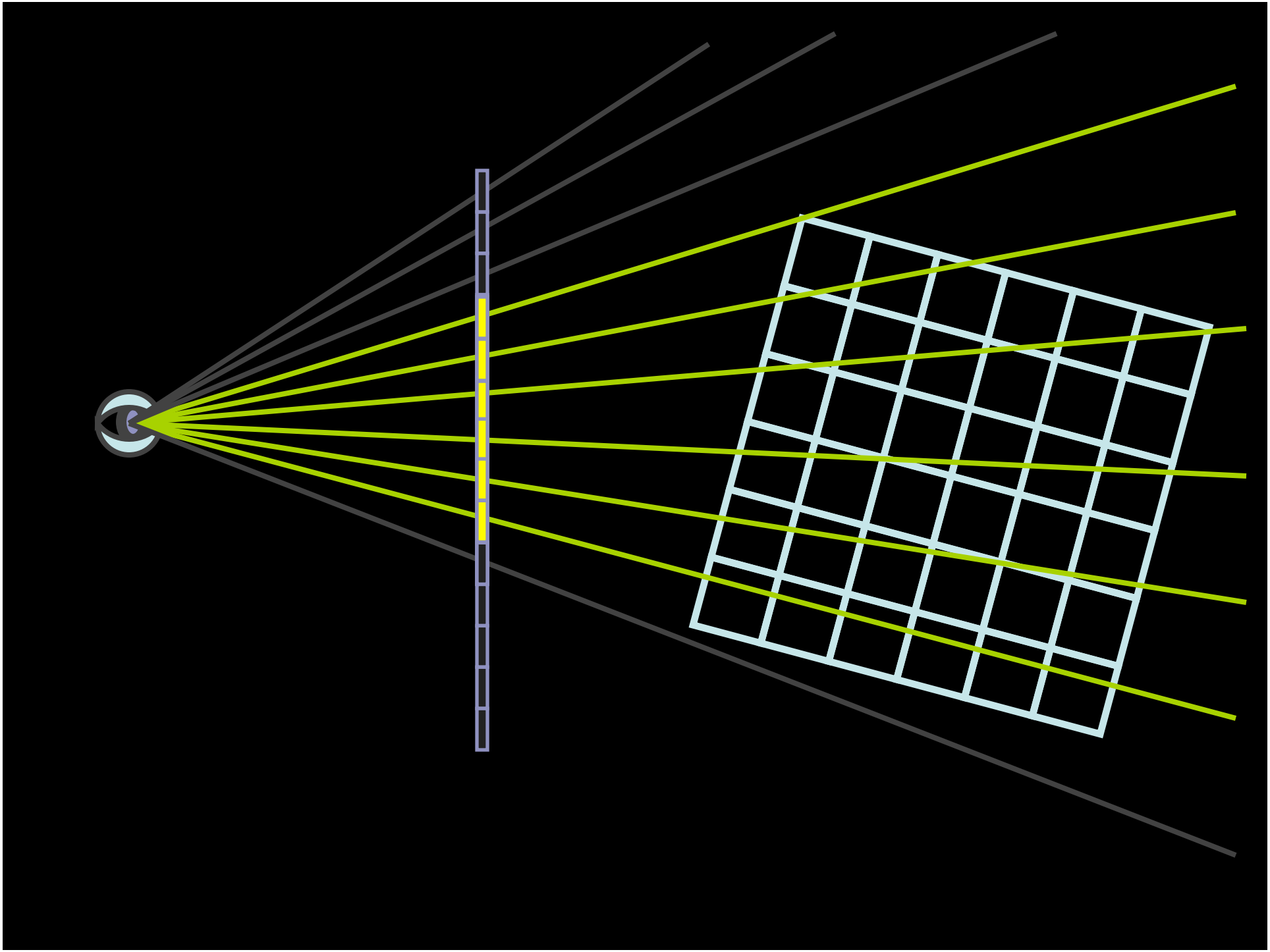
image space algorithm

object space algorithm

image space

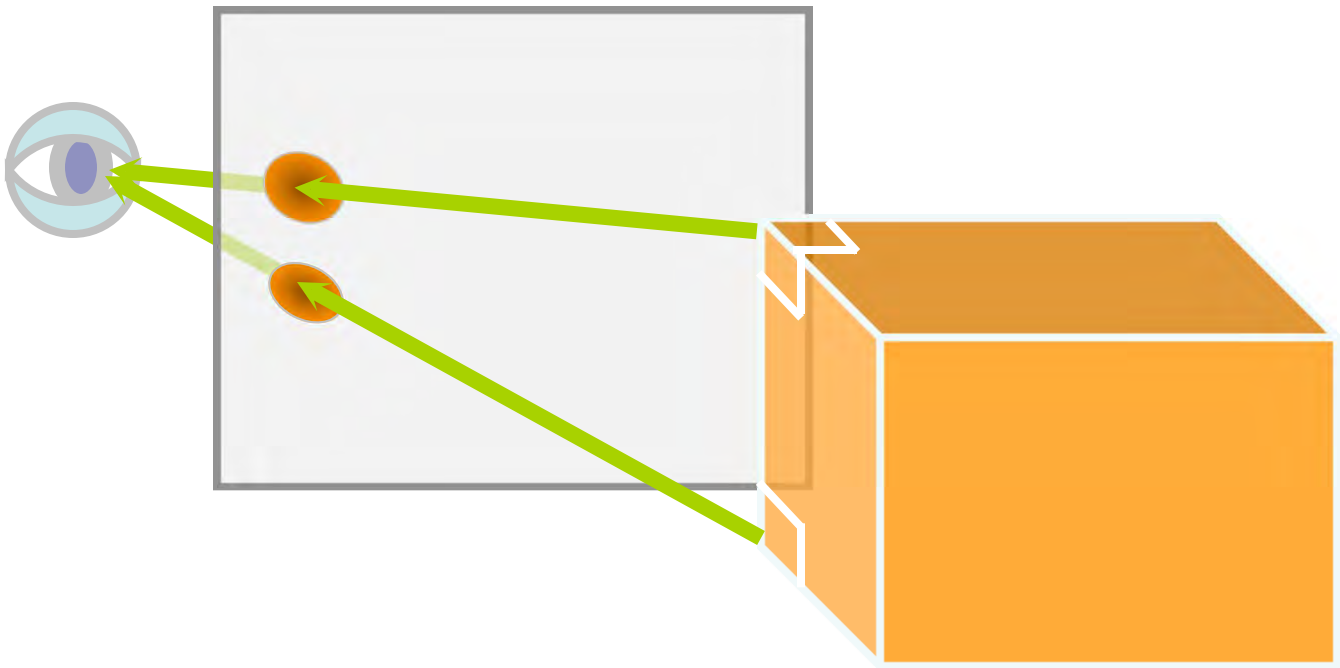
feed backward: ray casting

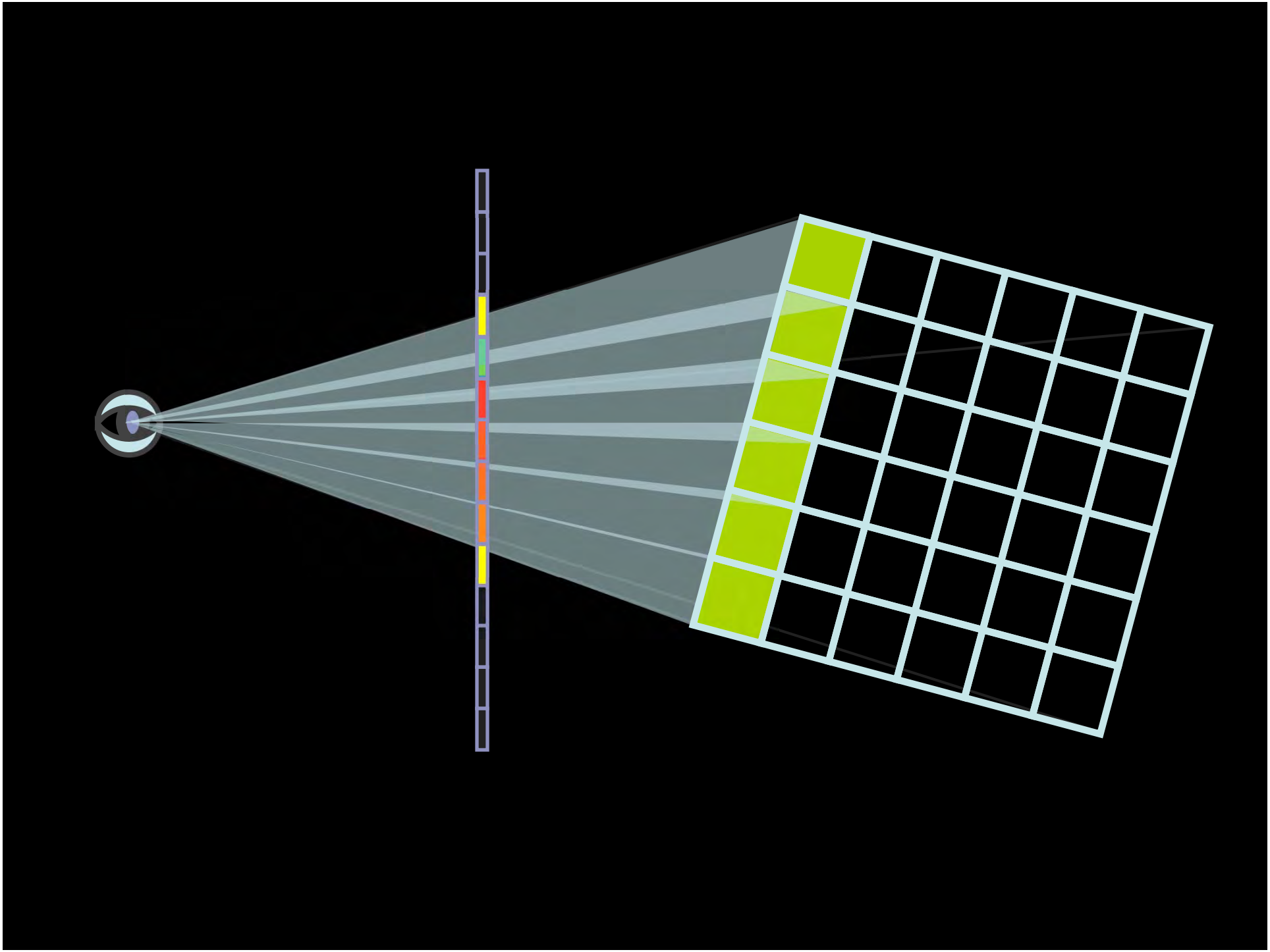




object space

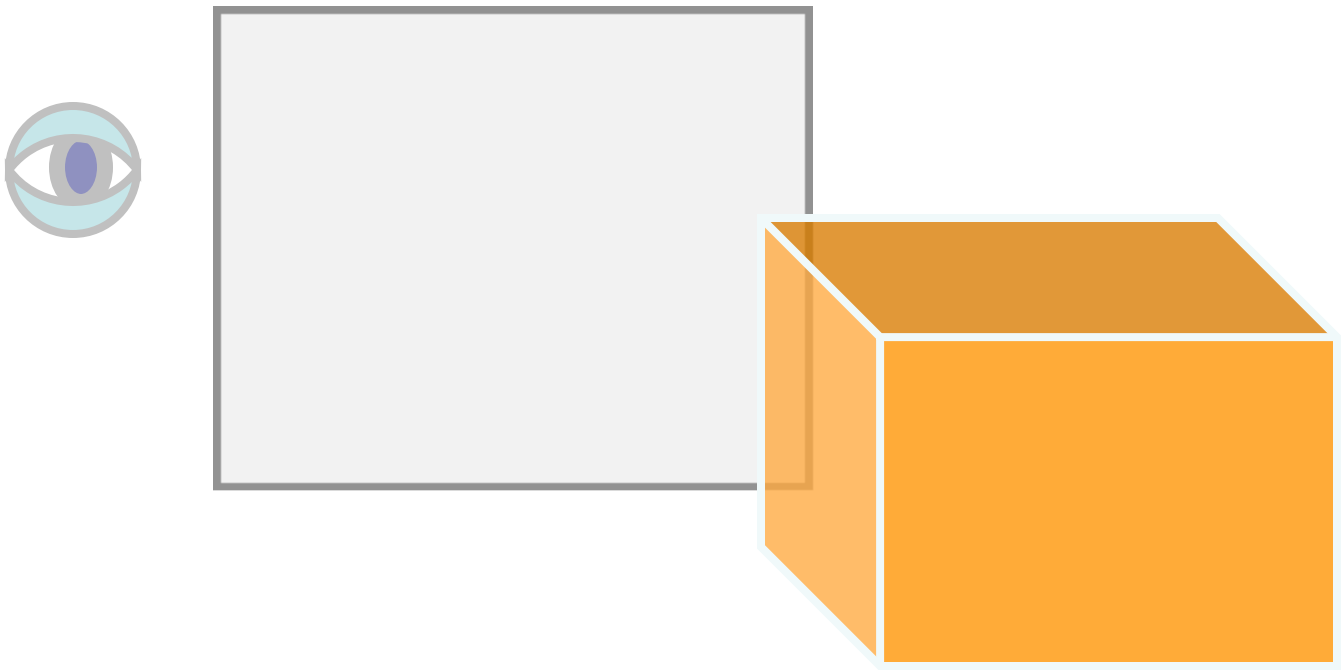
feed forward: splatting

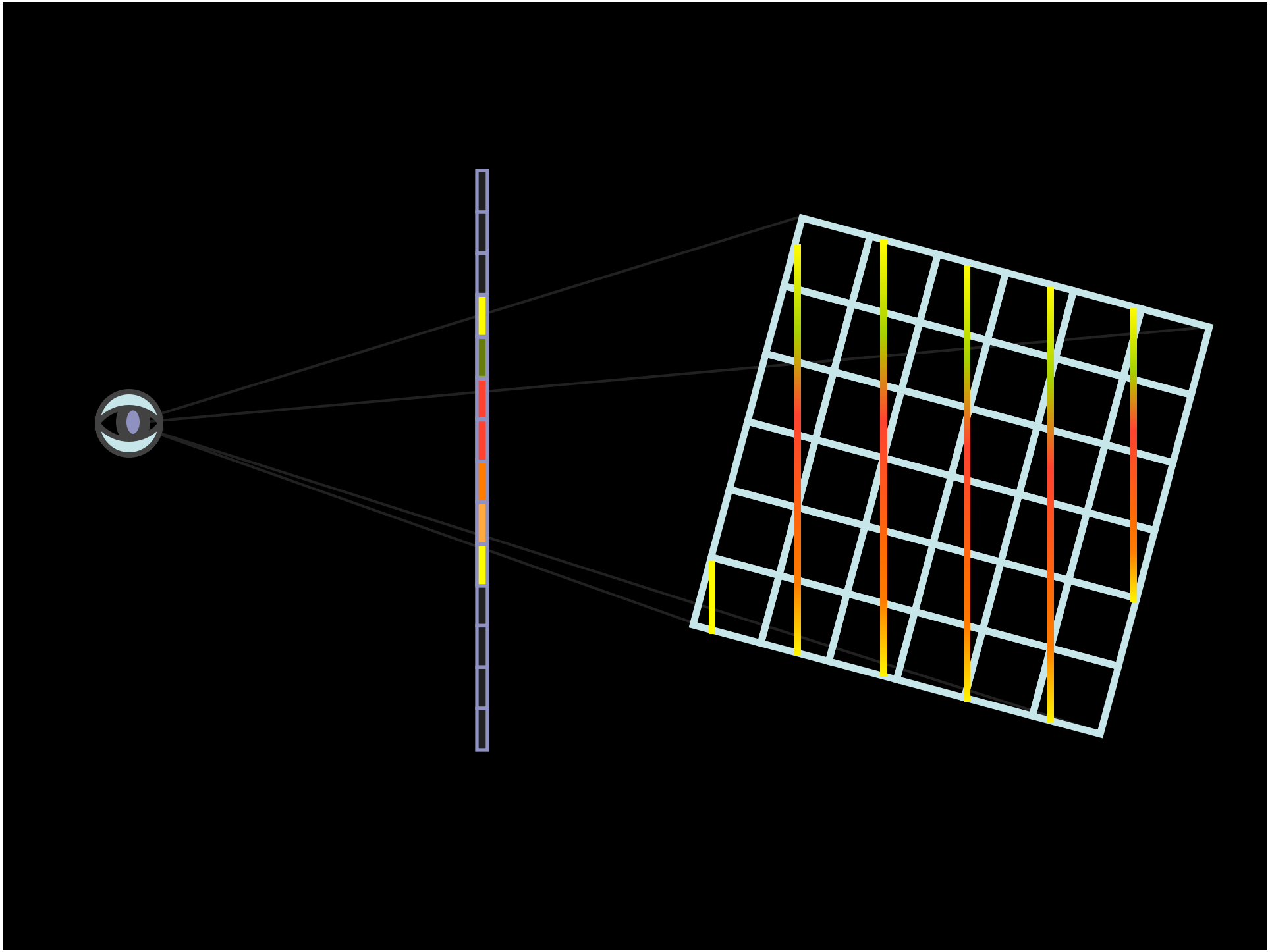




using *gpu*

triangulate slices: 3d textures

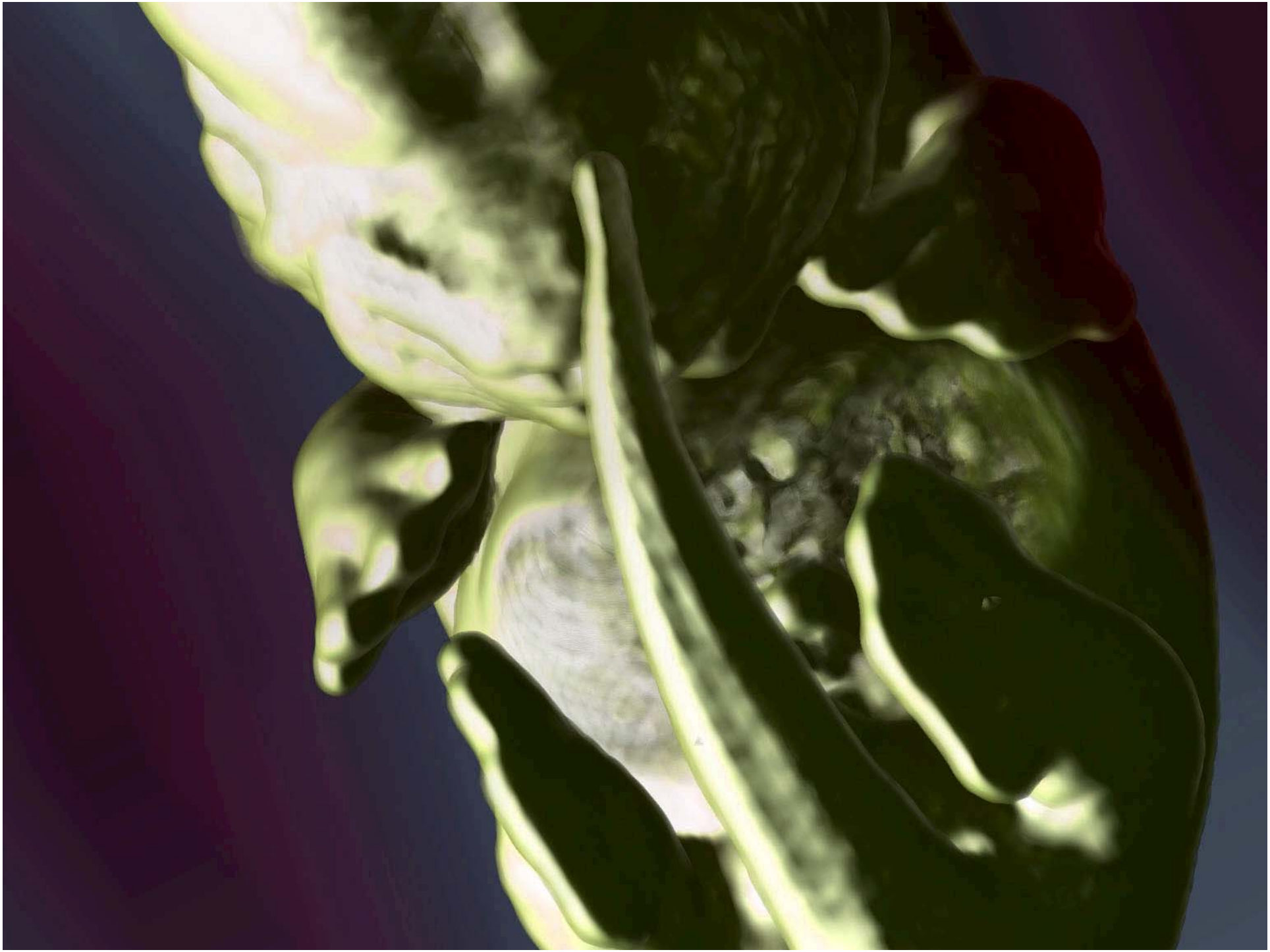




mouse ct scan:

PARAVIEW

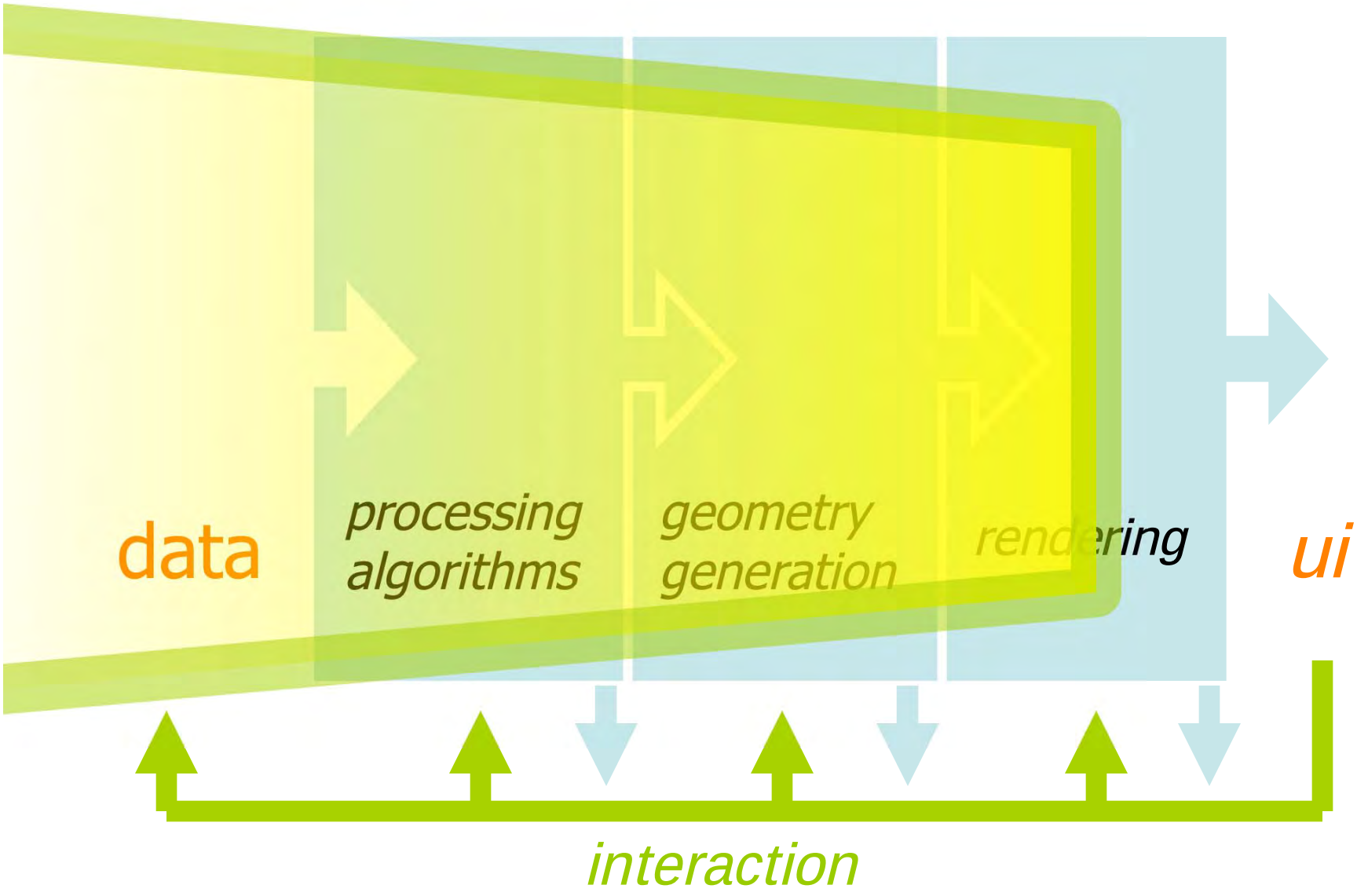
quick demo

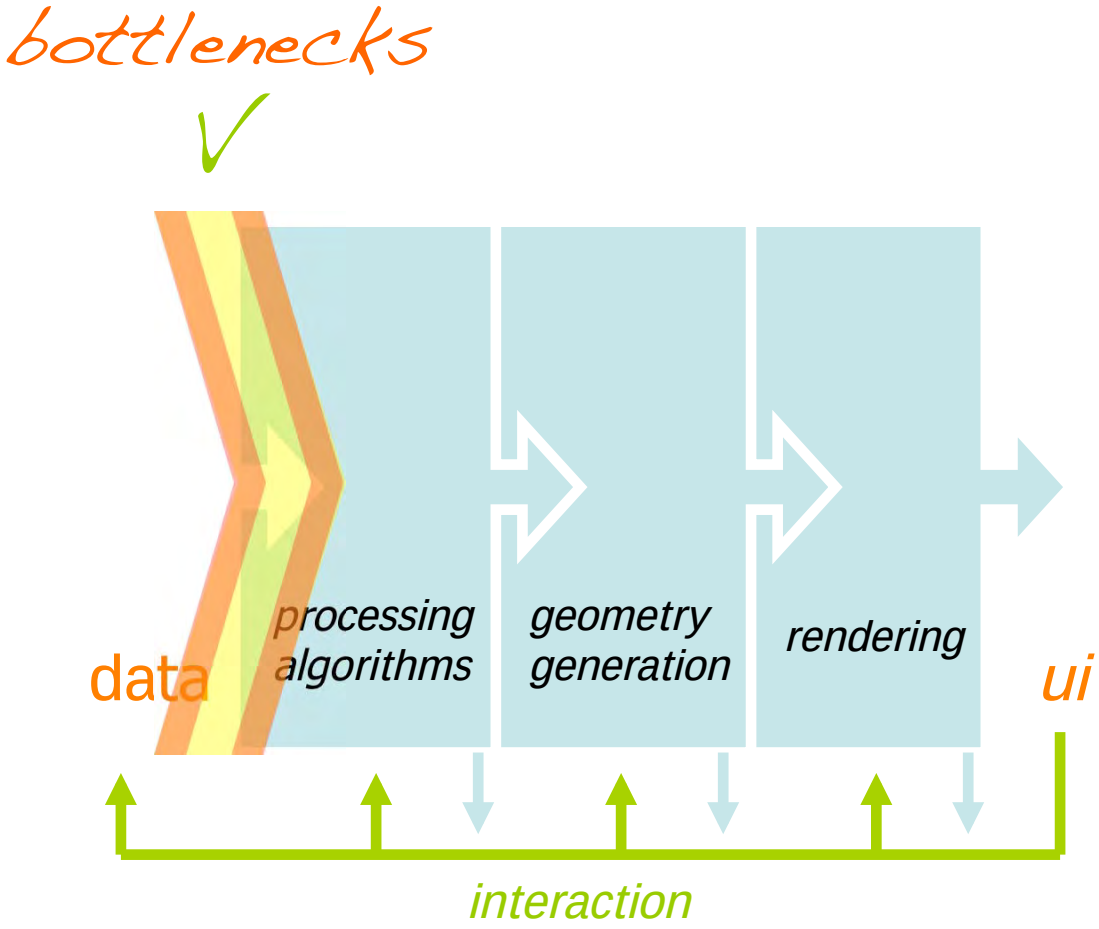


CLOSER LOOK:

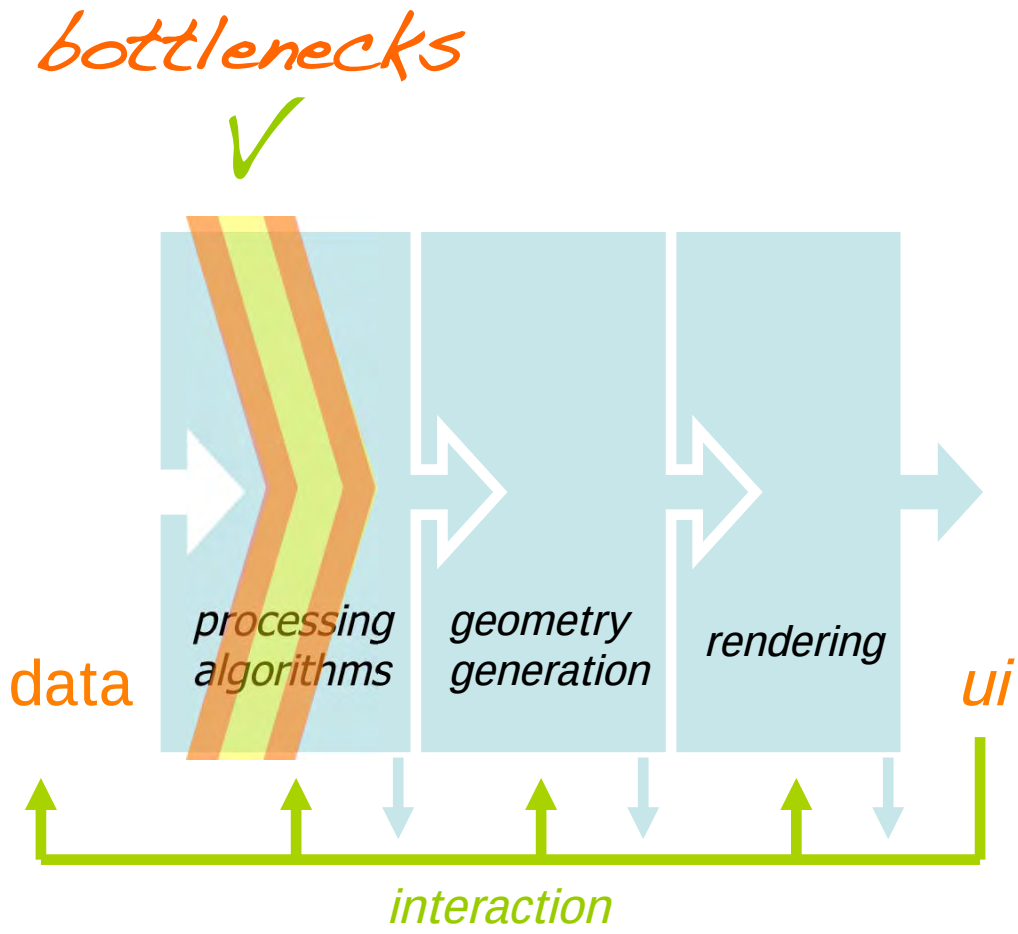
BOTTLENECKS

usual visualization “engine”



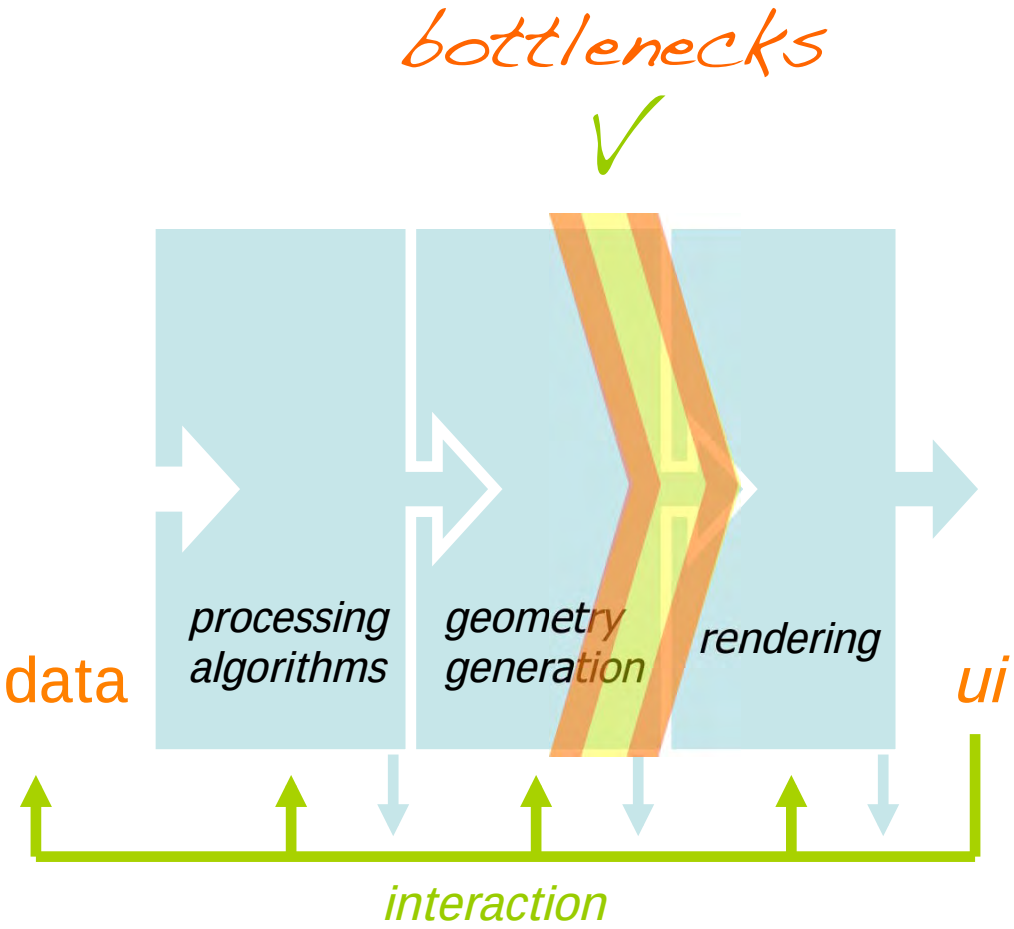


- data size
- data format
  - *xml*

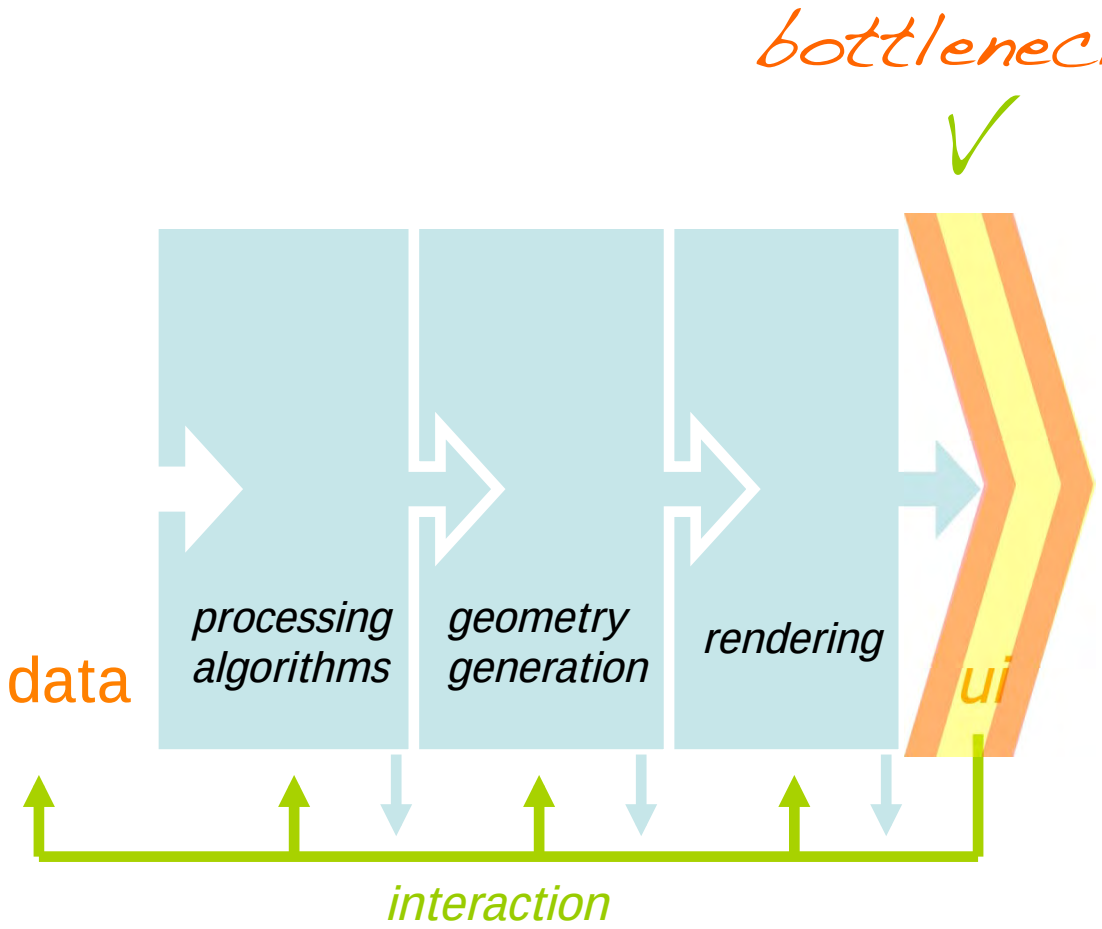


computation

power



- number of records
- number of triangles
  (base rendering units)
- number voxels



- knowledge level of "end user"
- complexity of data base

bottlenecks: *addressing the problem*

lod vs parallelism

addressing throughput bottlenecks

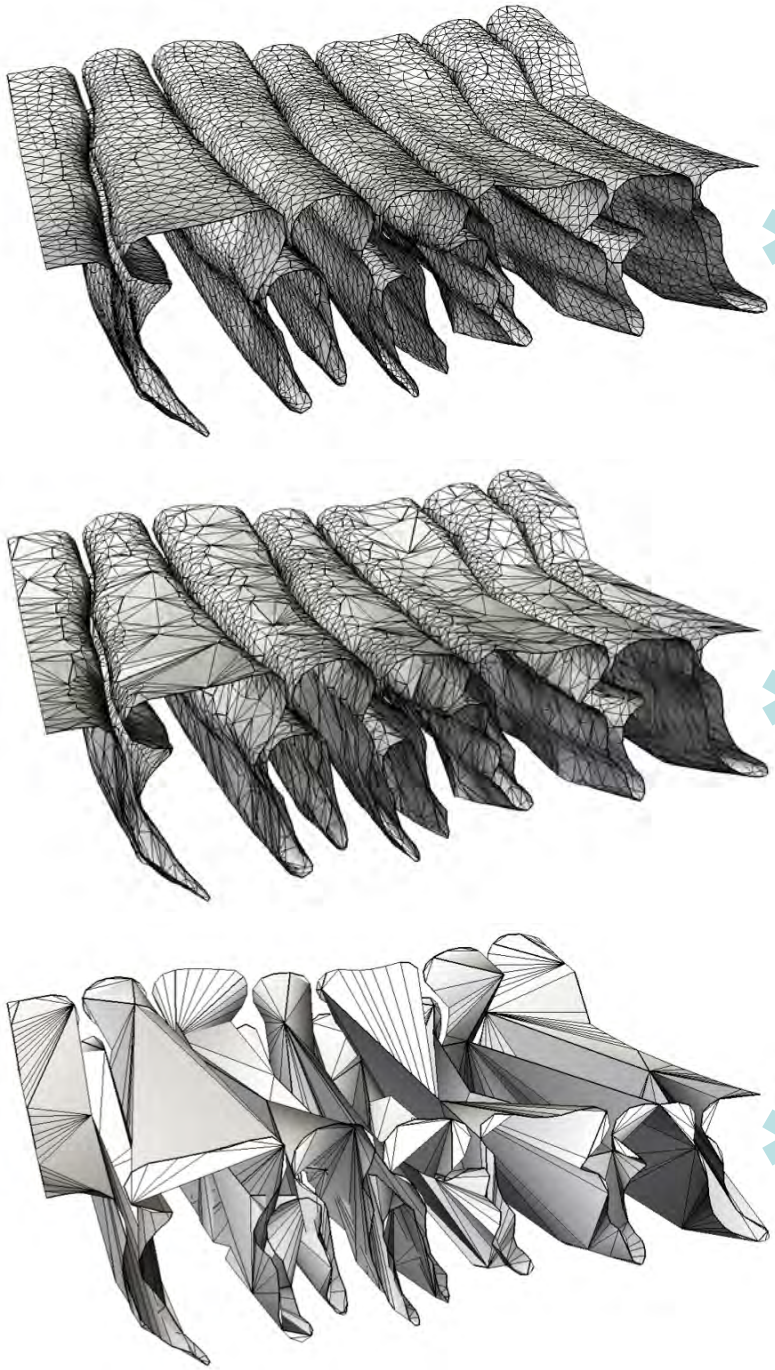
level of detail (LOD)

* requires pre-processing
* requires larger storage *(original+...)*

parallel processing/rendering

* requires a parallel system
* increases sw complexity
* less likely to be "portable"

lod: decimation



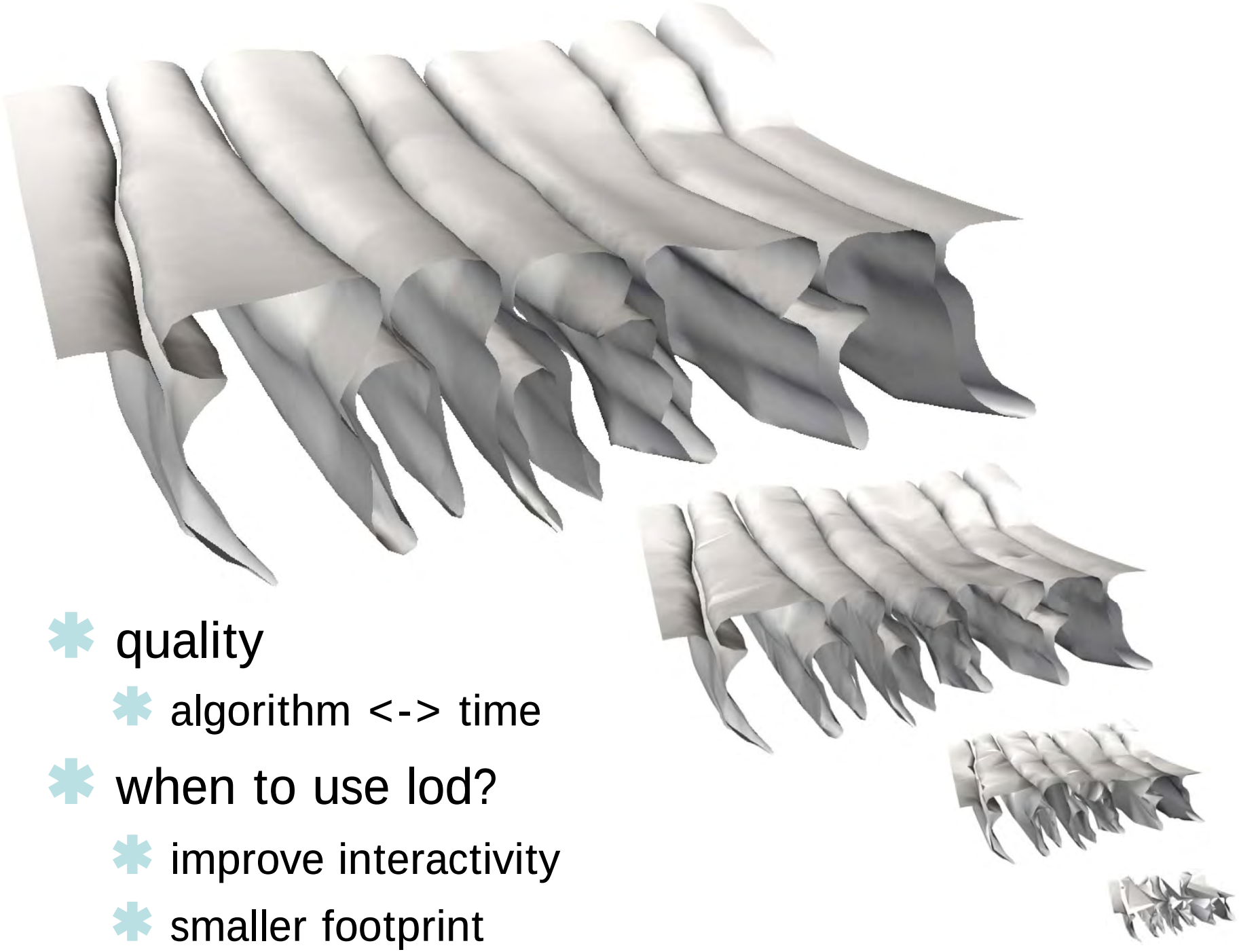
25365

12681

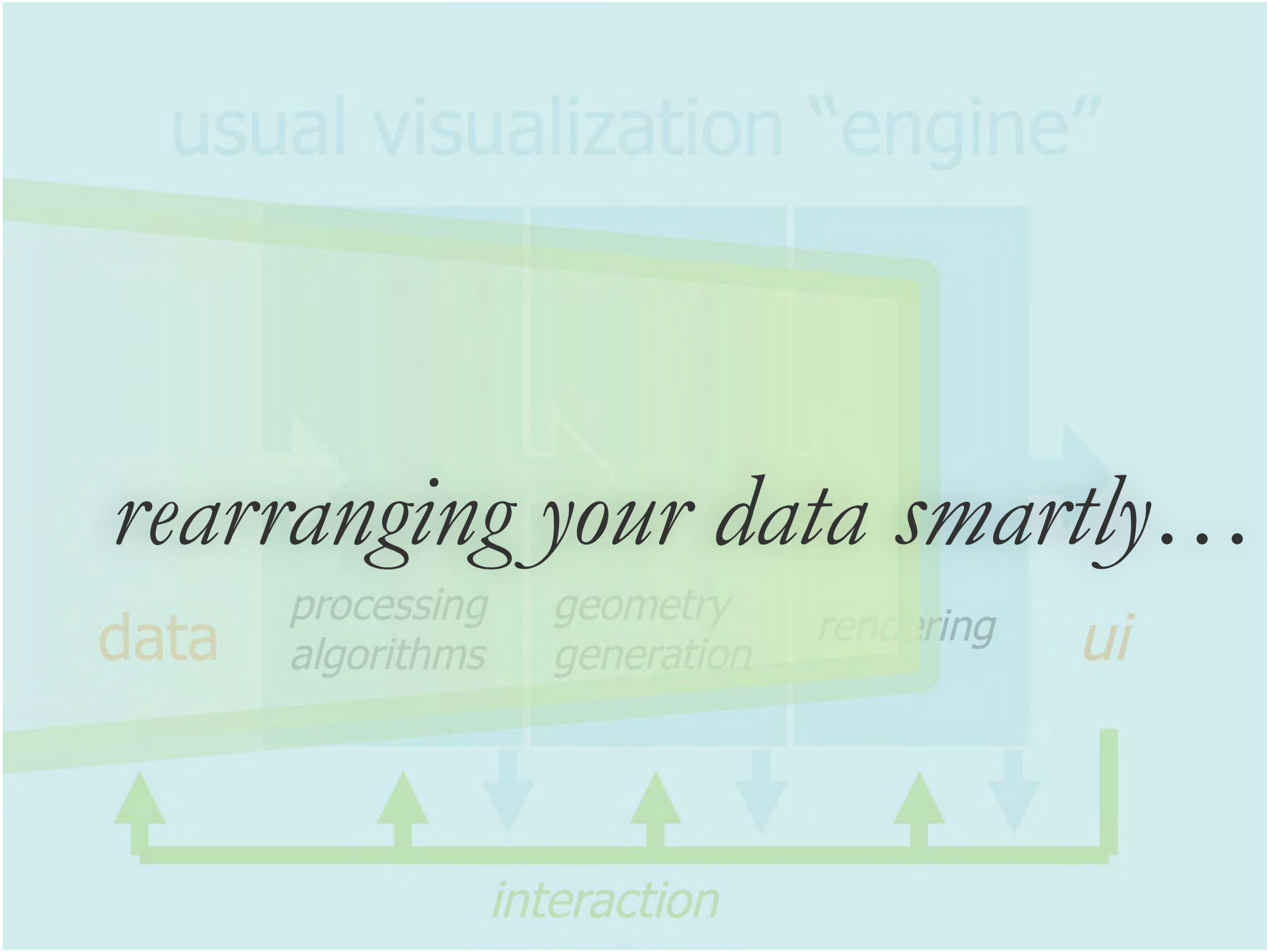
50%

1011

4%

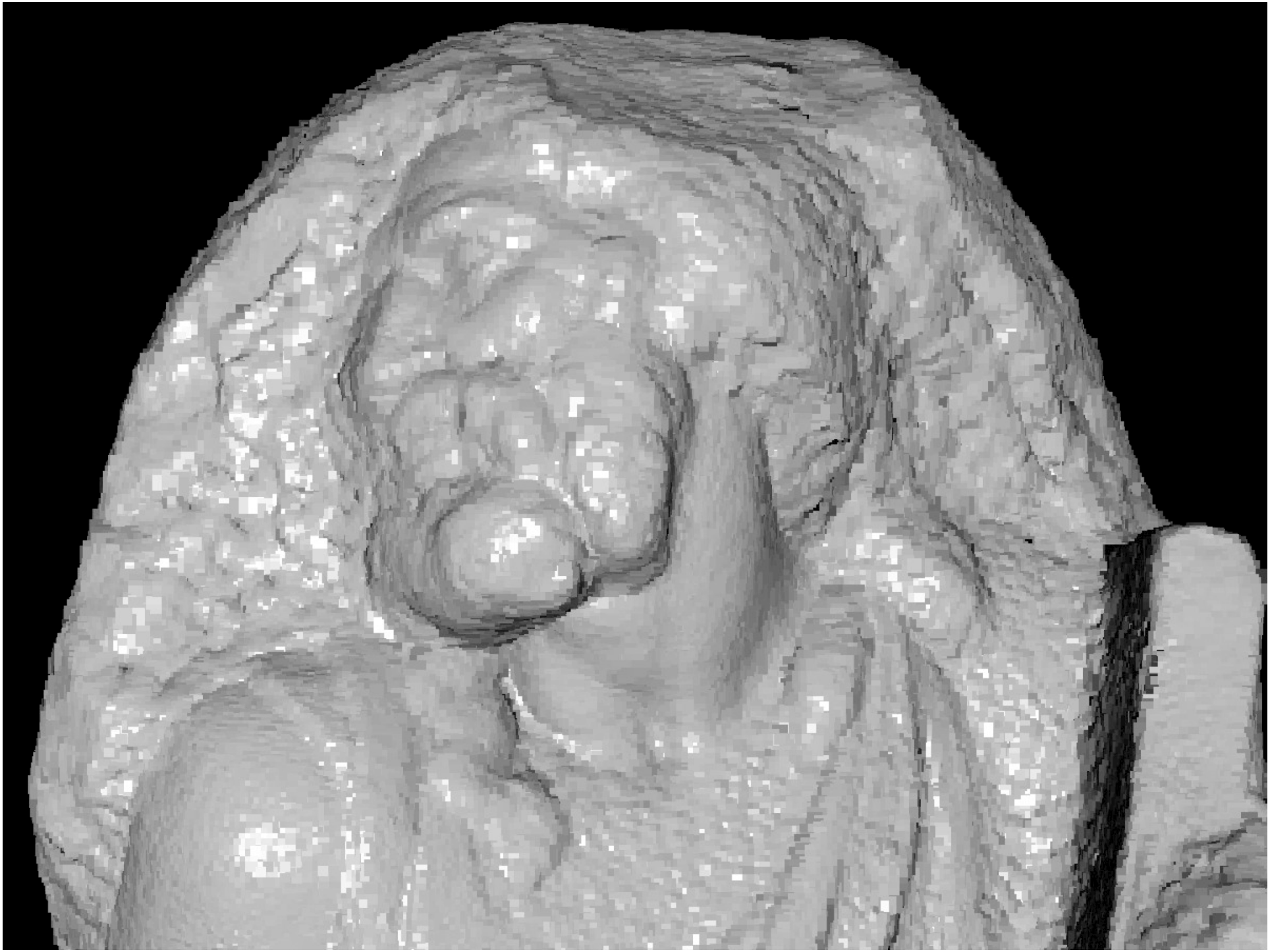


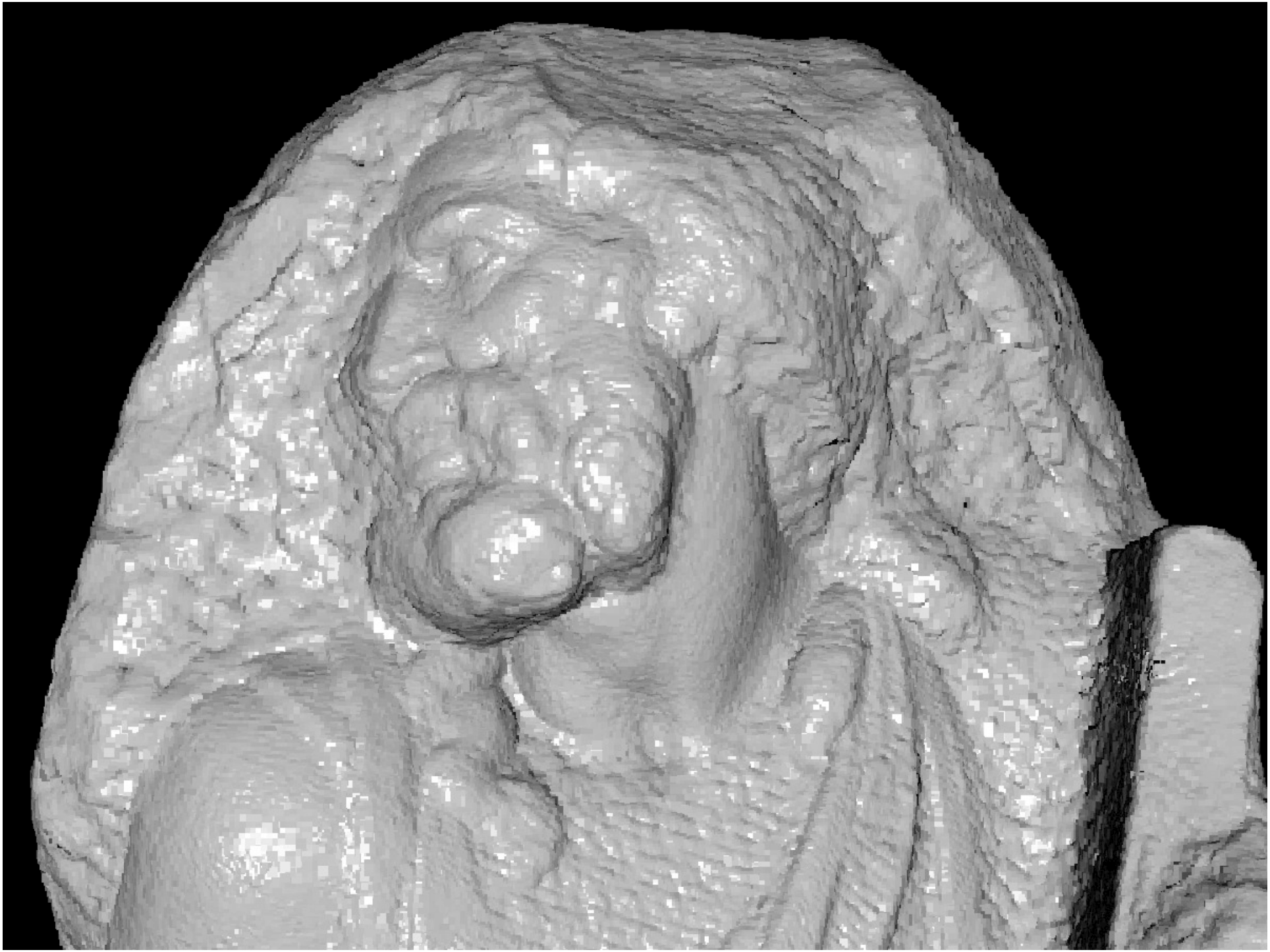
- quality
  - algorithm <-> time
- when to use lod?
  - improve interactivity
  - smaller footprint

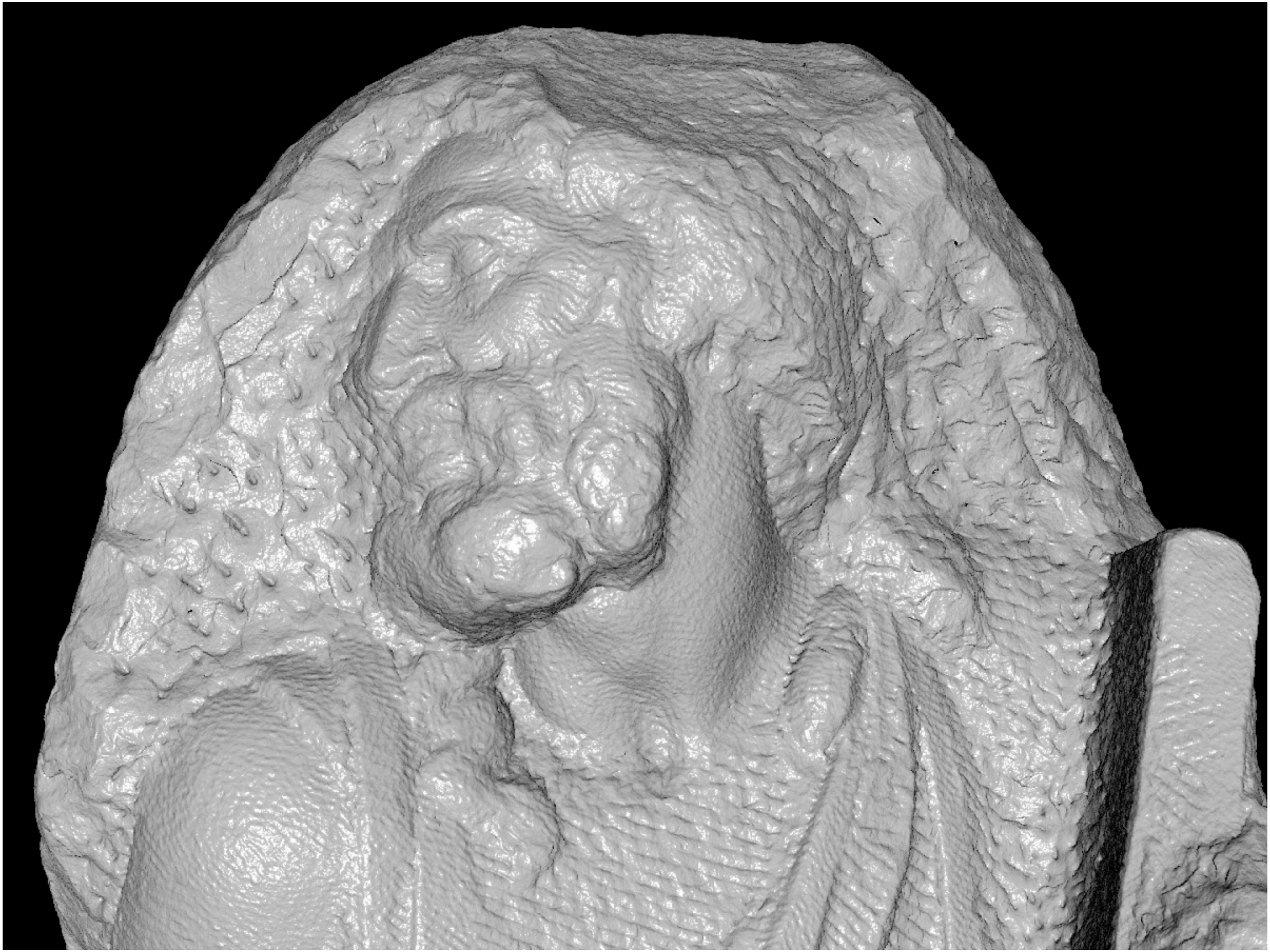


rearranging your data smartly…

stanford: qsplat

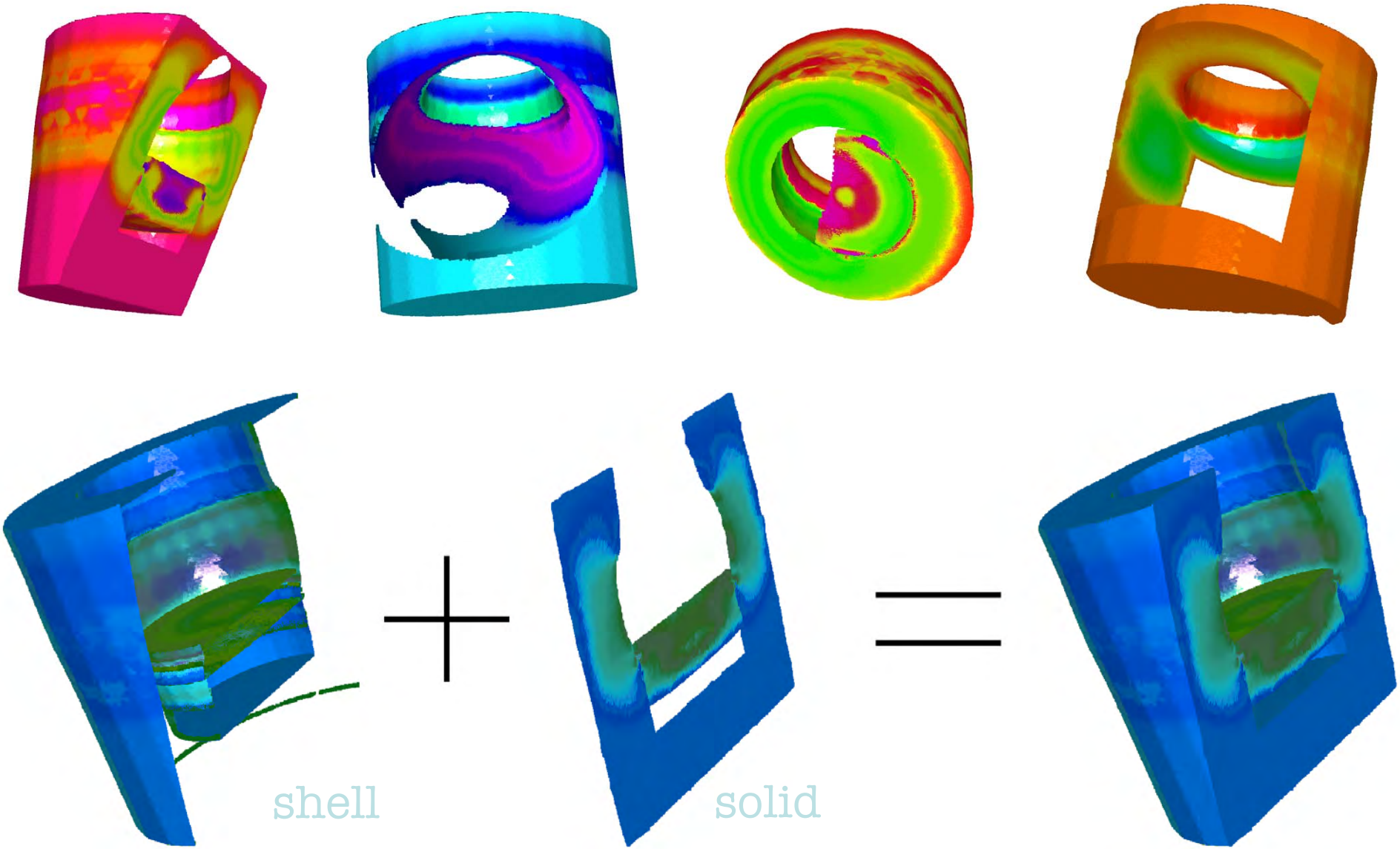






extending to volumetric data: tetsplat

exploring the shock-test cannister



bottlenecks: *addressing the problem*

lod vs parallelism

addressing throughput bottlenecks

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* requires larger storage

parallel processing/rendering

* requires a parallel system
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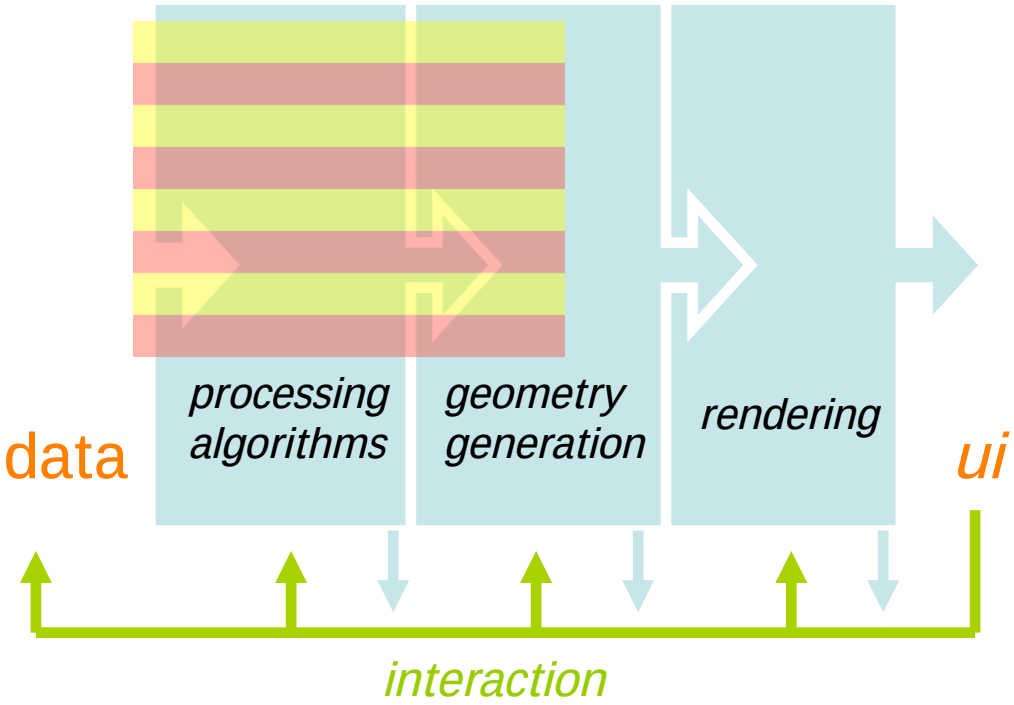
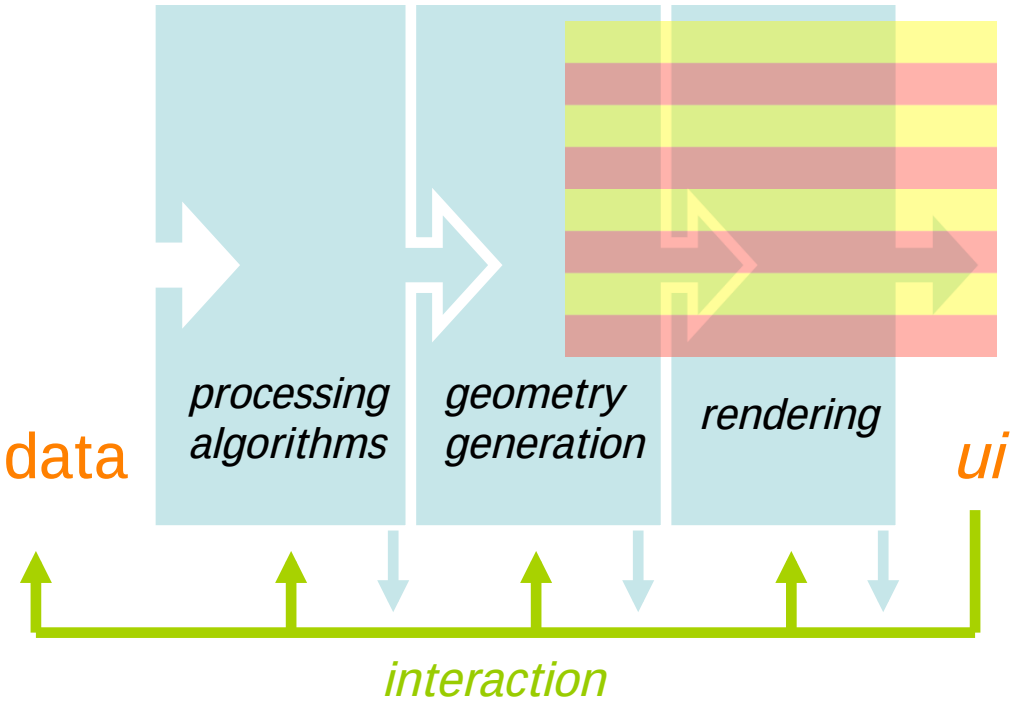
>> parallel viz >>

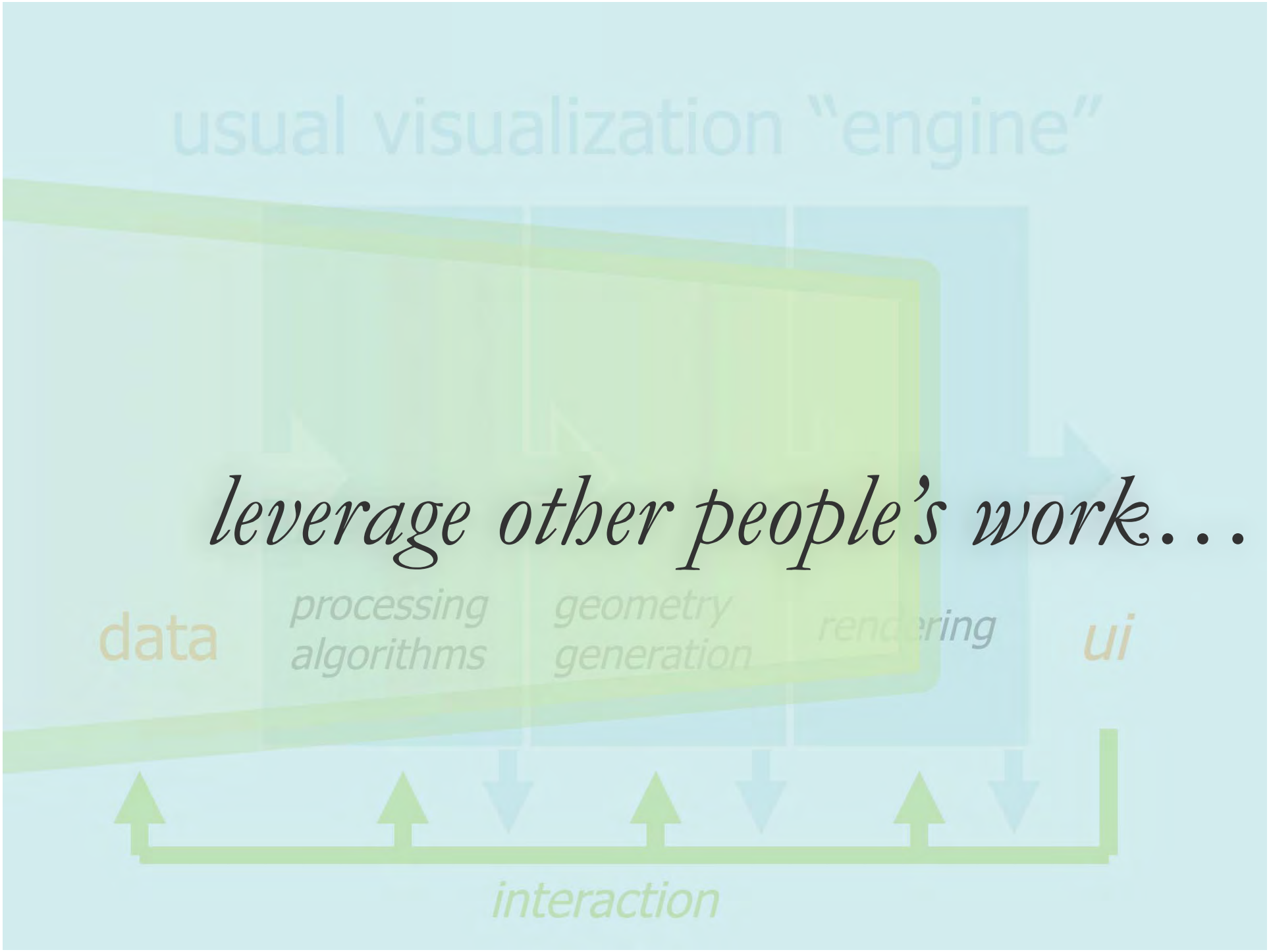


- + cpus
- data?
- seams?

>> parallel viz >>



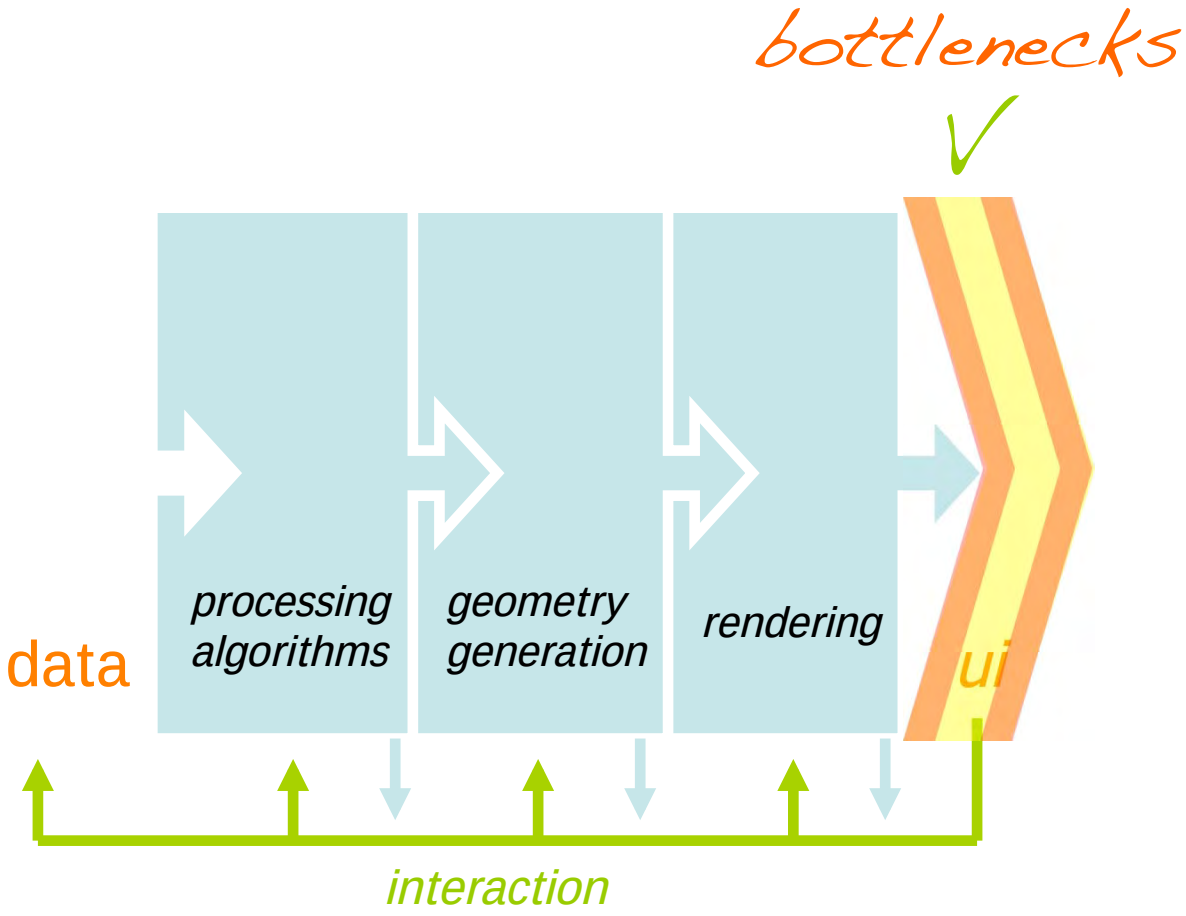
- + gpus
- screen?
- space?



leverage other people's work…

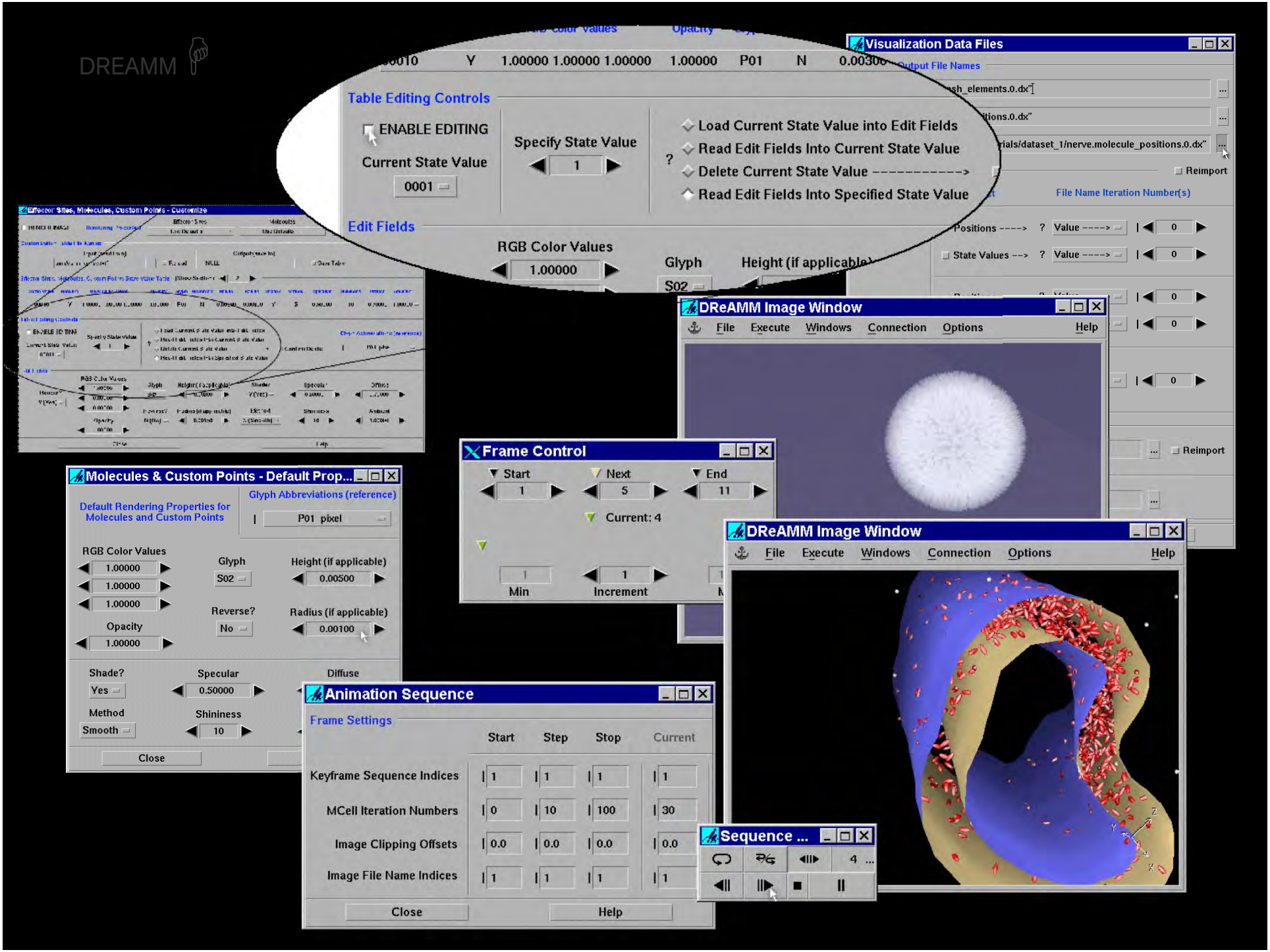
bottlenecks: *addressing the problem*

ui

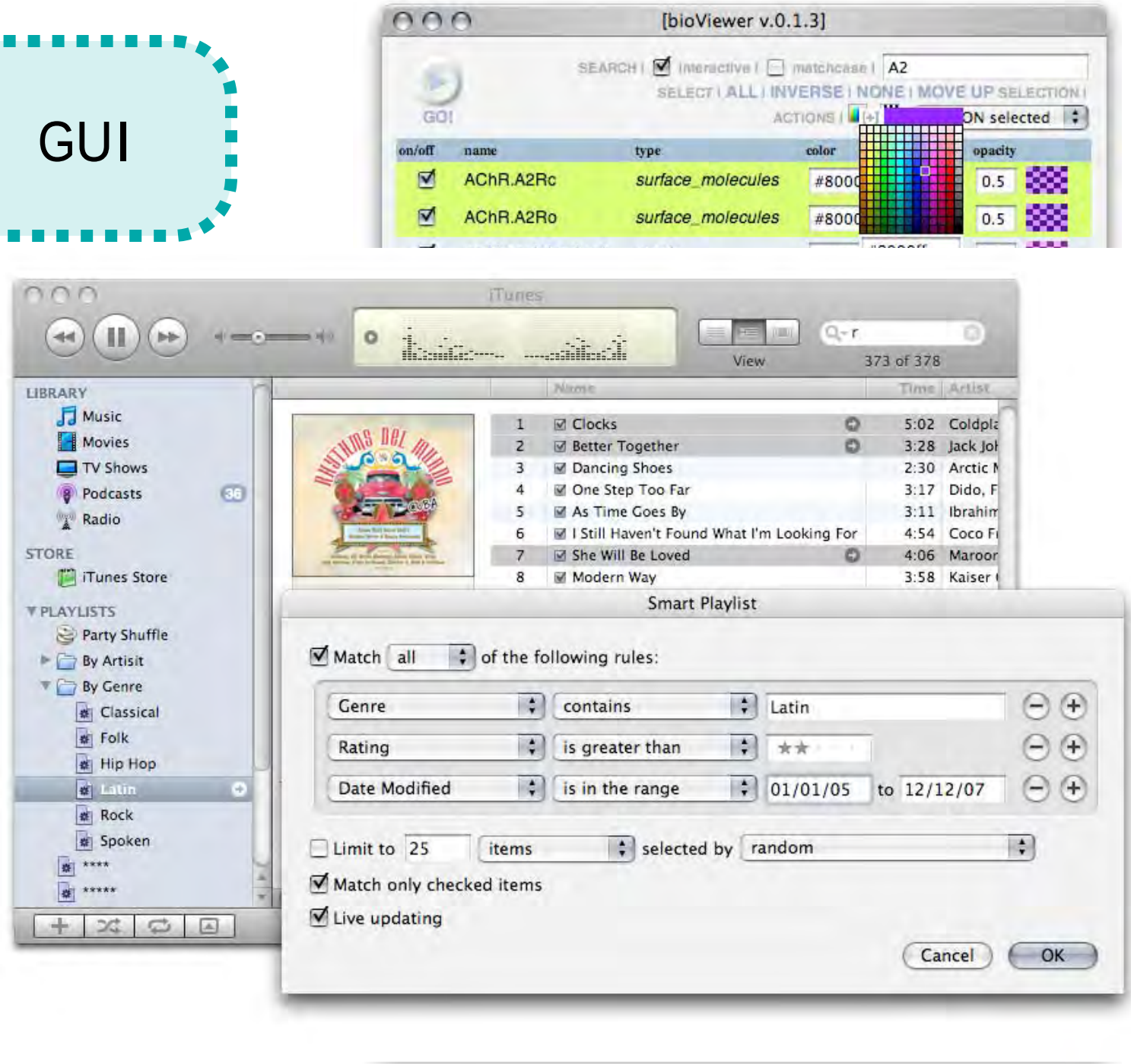

bottlenecks
data
processing algorithms
geometry generation
rendering
ui
interaction

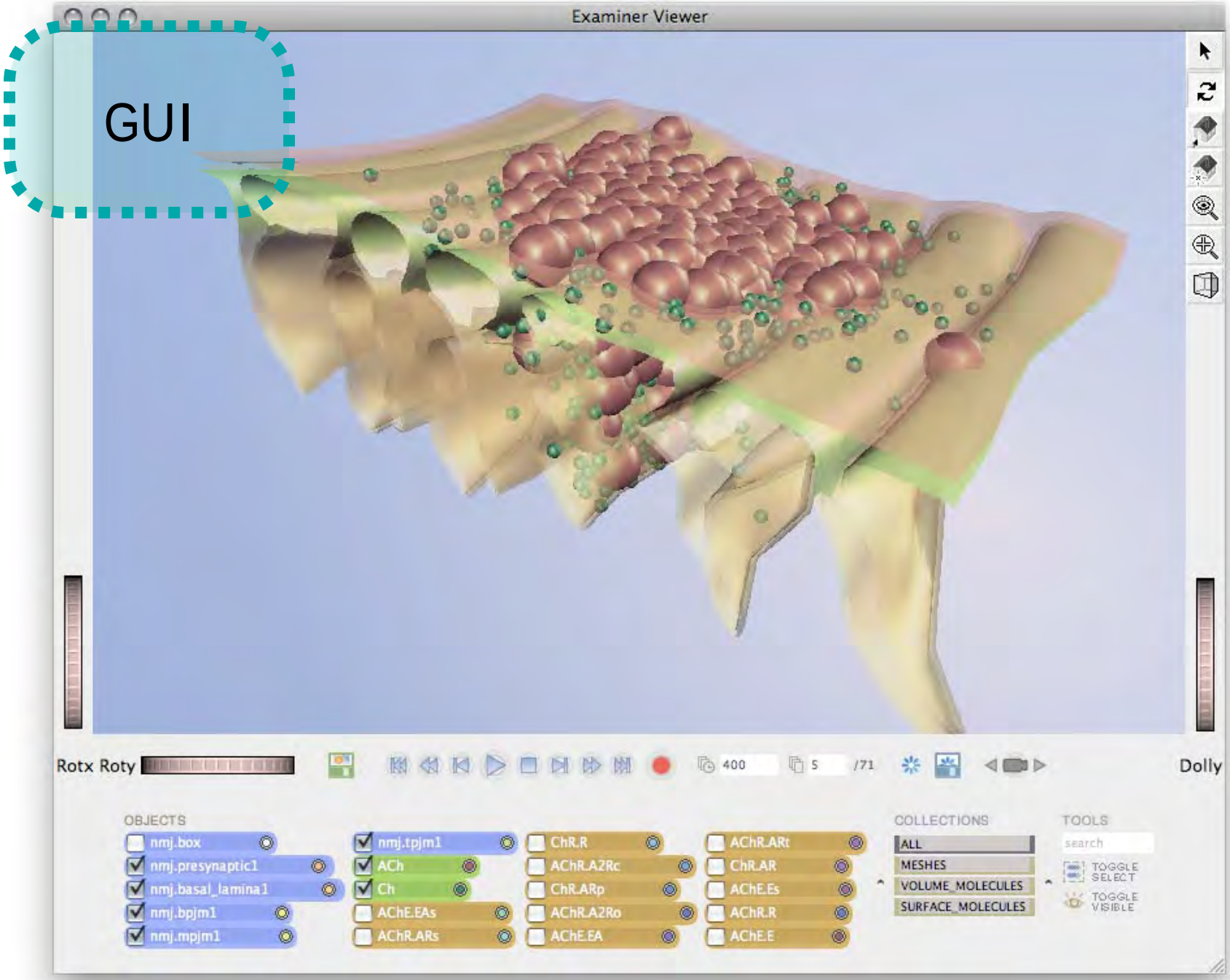
case study: *gui design*

mcell viewer

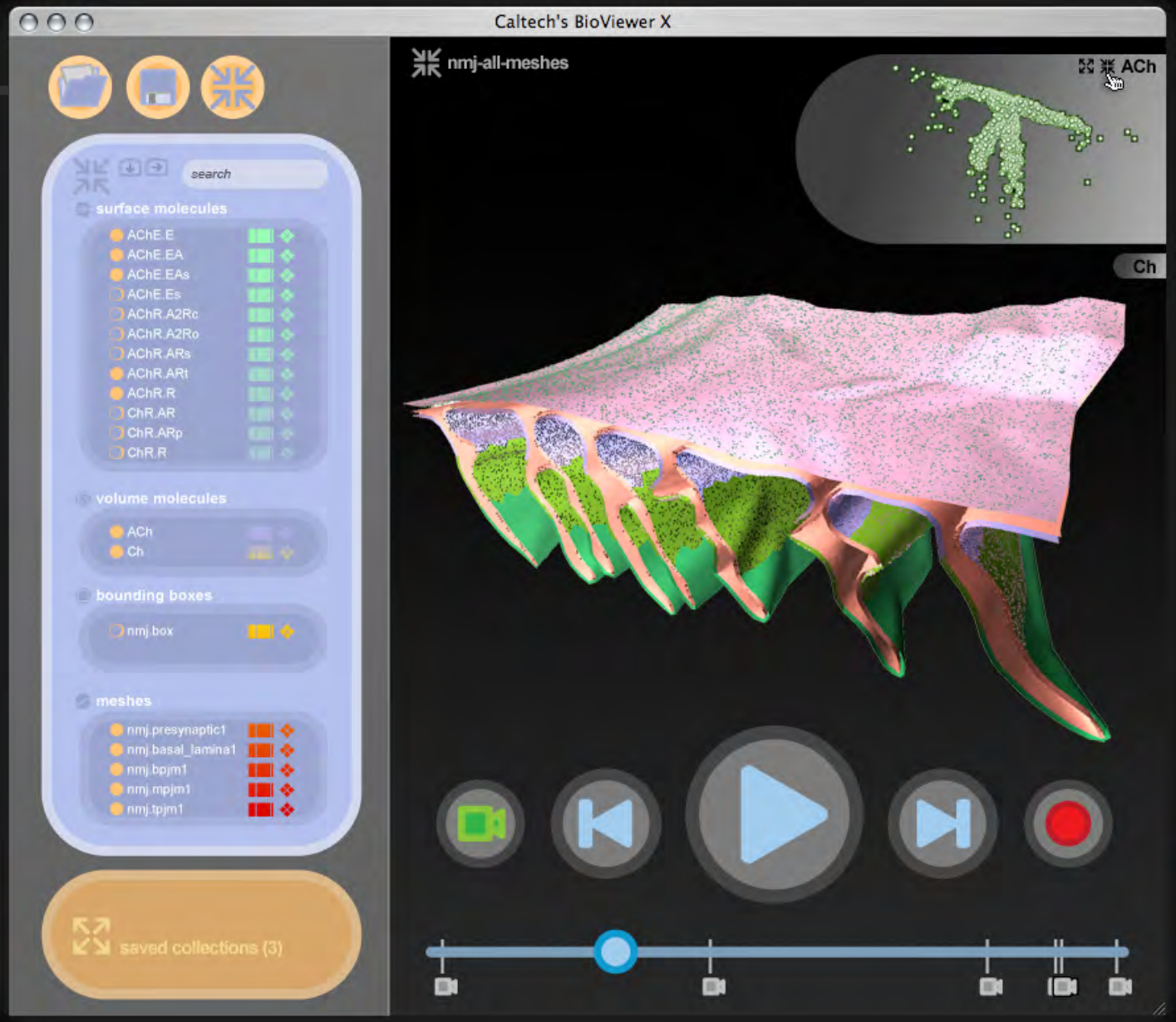

DREAMM
Table Editing Controls
ENABLE EDITING
Current State Value
0001
Specify State Value
1
Load Current State Value into Edit Fields
Read Edit Fields Into Current State Value
Delete Current State Value
Read Edit Fields Into Specified State Value
Edit Fields
RGB Color Values
1.00000
Glyph
S02
Height (if applicable)
Visualization Data Files
Output File Names
File Name Iteration Number(s)
Reimport
DReAMM Image Window
File
Execute
Windows
Connection
Options
Help
Frame Control
Start
Next
End
Current: 4
Min
Increment
Molecules & Custom Points - Default Prop...
Default Rendering Properties for Molecules and Custom Points
Glyph Abbreviations (reference)
P01 pixel
Opacity
Reverse?
No
Radius (if applicable)
0.00100
Shade?
Yes
Specular
0.50000
Diffuse
Method
Smooth
Shininess
10
Close
Animation Sequence
Frame Settings
Start
Step
Stop
Current
Keyframe Sequence Indices
MCell Iteration Numbers
Image Clipping Offsets
Image File Name Indices
Close
Help
Sequence ...

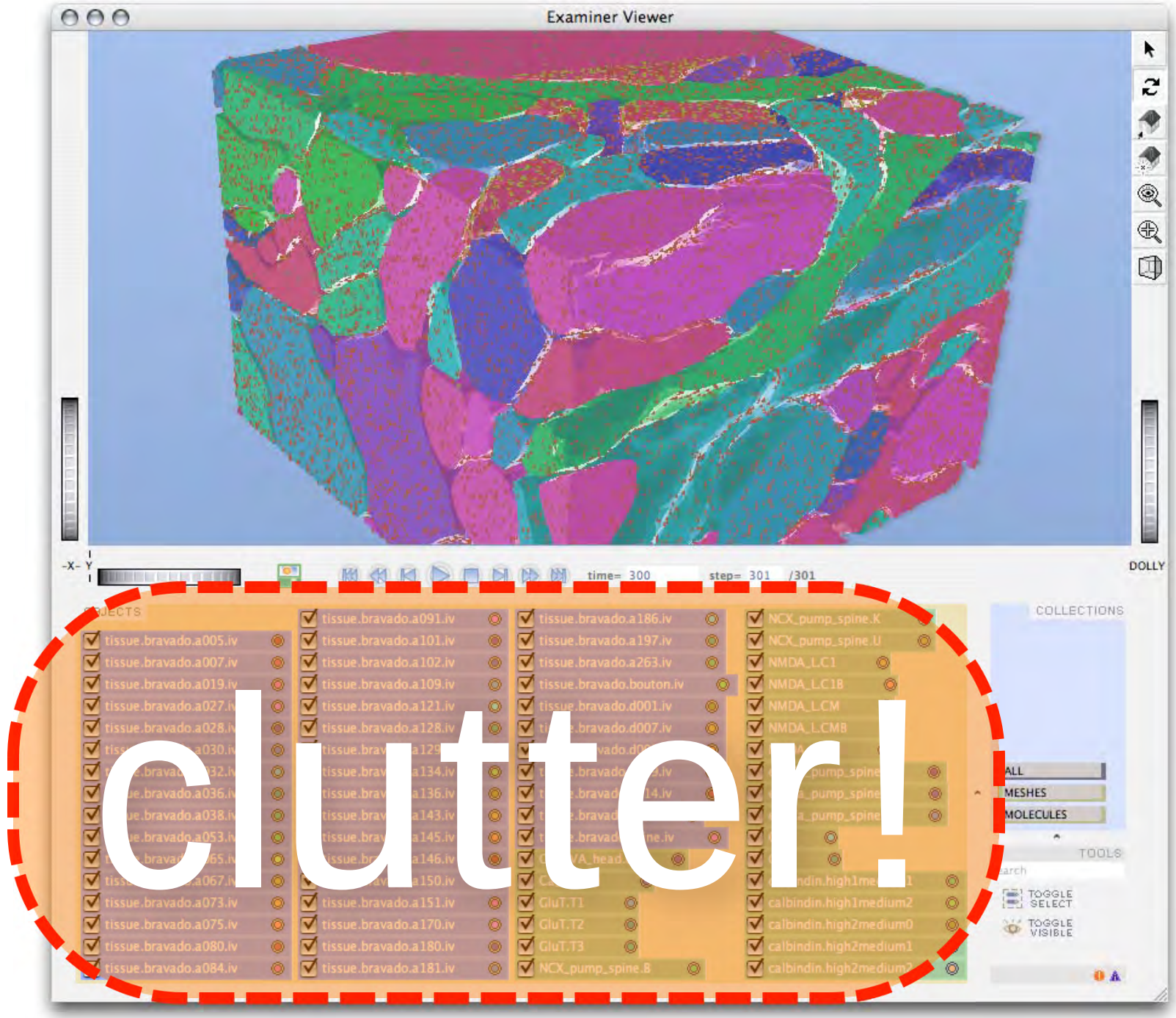
GUI

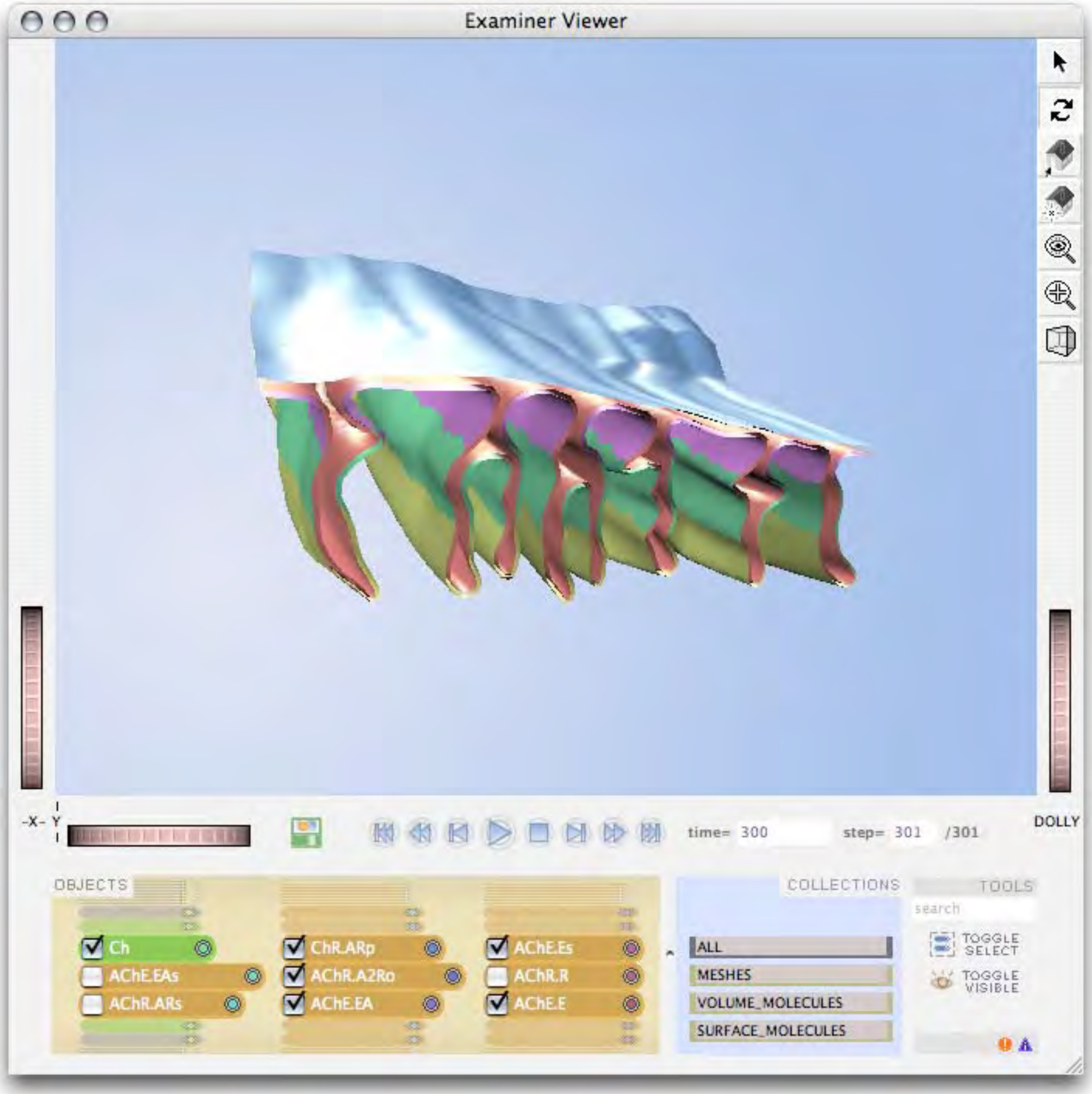

[bioViewer v.0.1.3]
SEARCH | Interactive | matchcase | A2
SELECT | ALL | INVERSE | NONE | MOVE UP SELECTION |
GO!
ACTIONS |
ON selected
on/off
name
type
color
opacity
AChR.A2Rc
surface_molecules
0.5
AChR.A2Ro
surface_molecules
0.5
iTunes
View
373 of 378
LIBRARY
Music
Movies
TV Shows
Podcasts
Radio
STORE
iTunes Store
PLAYLISTS
Party Shuffle
By Artisit
By Genre
Classical
Folk
Hip Hop
Latin
Rock
Spoken
Name
Time
Artist
1 Clocks 5:02
2 Better Together 3:28
3 Dancing Shoes 2:30
4 One Step Too Far 3:17
5 As Time Goes By 3:11
6 I Still Haven't Found What I'm Looking For 4:54
7 She Will Be Loved 4:06
8 Modern Way 3:58
Smart Playlist
Match all of the following rules:
Genre contains Latin
Rating is greater than
Date Modified is in the range 01/01/05 to 12/12/07
Limit to 25 items selected by random
Match only checked items
Live updating
Cancel
OK


GUI
Examiner Viewer
Rotx Roty
400
5
/71
Dolly
OBJECTS
nmj.box
nmj.presynaptic1
nmj.basal_lamina1
nmj.bpjm1
nmj.mpjm1
nmj.tpjm1
ACh
Ch
AChE.EAs
AChR.ARs
ChR.R
AChR.A2Rc
ChR.ARp
AChR.A2Ro
AChE.EA
AChR.ARt
ChR.AR
AChE.Es
AChR.R
AChE.E
COLLECTIONS
ALL
MESHES
VOLUME_MOLECULES
SURFACE_MOLECULES
TOOLS
search
TOGGLE SELECT
TOGGLE VISIBLE

ultimate tool mockup




Examiner Viewer
time= 300
step= 301 /301
DOLLY
COLLECTIONS
ALL
MESHES
MOLECULES
TOOLS
TOGGLE SELECT
TOGGLE VISIBLE
tissue.bravado.a005.iv
tissue.bravado.a007.iv
tissue.bravado.a019.iv
tissue.bravado.a027.iv
tissue.bravado.a028.iv
tissue.bravado.a073.iv
tissue.bravado.a075.iv
tissue.bravado.a080.iv
tissue.bravado.a084.iv
tissue.bravado.a091.iv
tissue.bravado.a101.iv
tissue.bravado.a102.iv
tissue.bravado.a109.iv
tissue.bravado.a121.iv
tissue.bravado.a128.iv
tissue.bravado.a151.iv
tissue.bravado.a170.iv
tissue.bravado.a180.iv
tissue.bravado.a181.iv
tissue.bravado.a186.iv
tissue.bravado.a197.iv
tissue.bravado.a263.iv
tissue.bravado.bouton.iv
GluT.T1
GluT.T2
GluT.T3
NCX_pump_spine.B
NCX_pump_spine.K
NCX_pump_spine.U
NMDA_LC1
NMDA_LC1B
NMDA_LCM
NMDA_LCMB
calbindin.high1medium2
calbindin.high2medium0
calbindin.high2medium1
clutter!

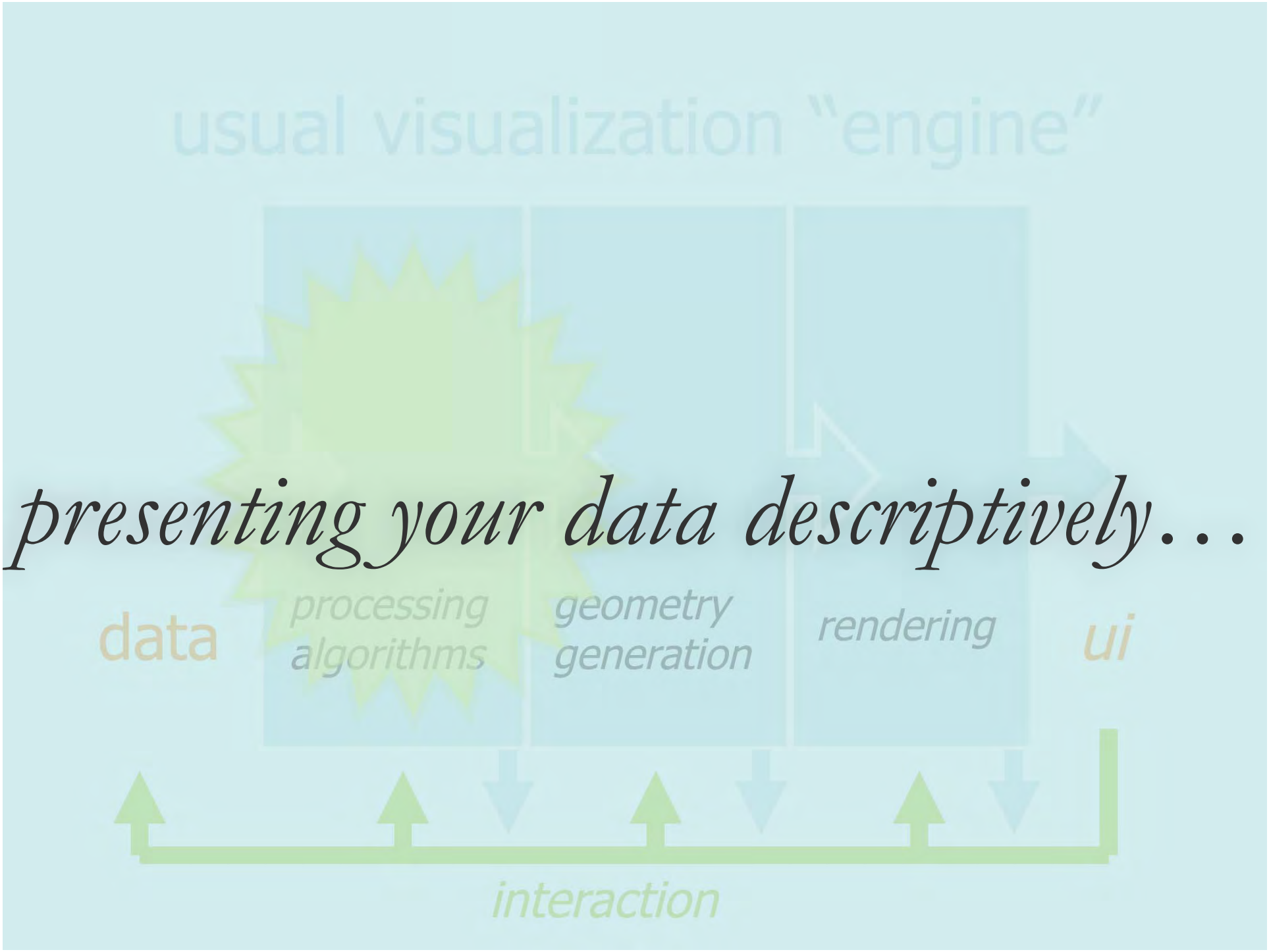

Examiner Viewer
-X-
Y
DOLLY
time= 300
step= 301
/301
OBJECTS
Ch
AChE.EAs
AChR.ARs
ChR.ARp
AChR.A2Ro
AChE.EA
AChE.Es
AChR.R
AChE.E
COLLECTIONS
ALL
MESHES
VOLUME_MOLECULES
SURFACE_MOLECULES
TOOLS
search
TOGGLE SELECT
TOGGLE VISIBLE

ibm: wordle


visualization
data
bottlenecks
based
engine
parallel
advanced
python
requires
computing
STRUCTURE
GEOMETRIC
abstract
research
caltech
center
system
lombeyda
vtk
DataArray
techniques
number
space
VTK
usual
santiago
representation
addressing
complexity
sample
volume
lab
rendering
viz
basic
binary
type
DATA
algorithm
level
look
format
PARAVIEW
ImageData
ascii
object
quick
toolkit
scalar
lod
xml
field
image
sketch
interactive
design
LOD
Name
topics
regular
algorithms
volumetric
voxels
maneesh
original+
Scalars
gui
tools
purpose
nmj
standard
Many
mapping
meshes
throughput
PointData
cellData
case
increases
preprocessing
ParaviewVTK
vs
study
agrawala
LLNL
interaction
looking
wiki
BIO
version
intuitive
Mollegro
grids
know
likely
c/c++
visualize
ray
java
base
EyesWWW
less
analysis
storage
VTKFile
larger
want
Java
raw
CellData
detail
portable
QT
VisitVTK
understandable
ProtovisWWW
TomCat
national
Wiki
feed
infovis
tcl/tk
EnSight
sw
Piece
processing/rendering
user
MCELL
pipeline
ModrianR

ibm: wordle

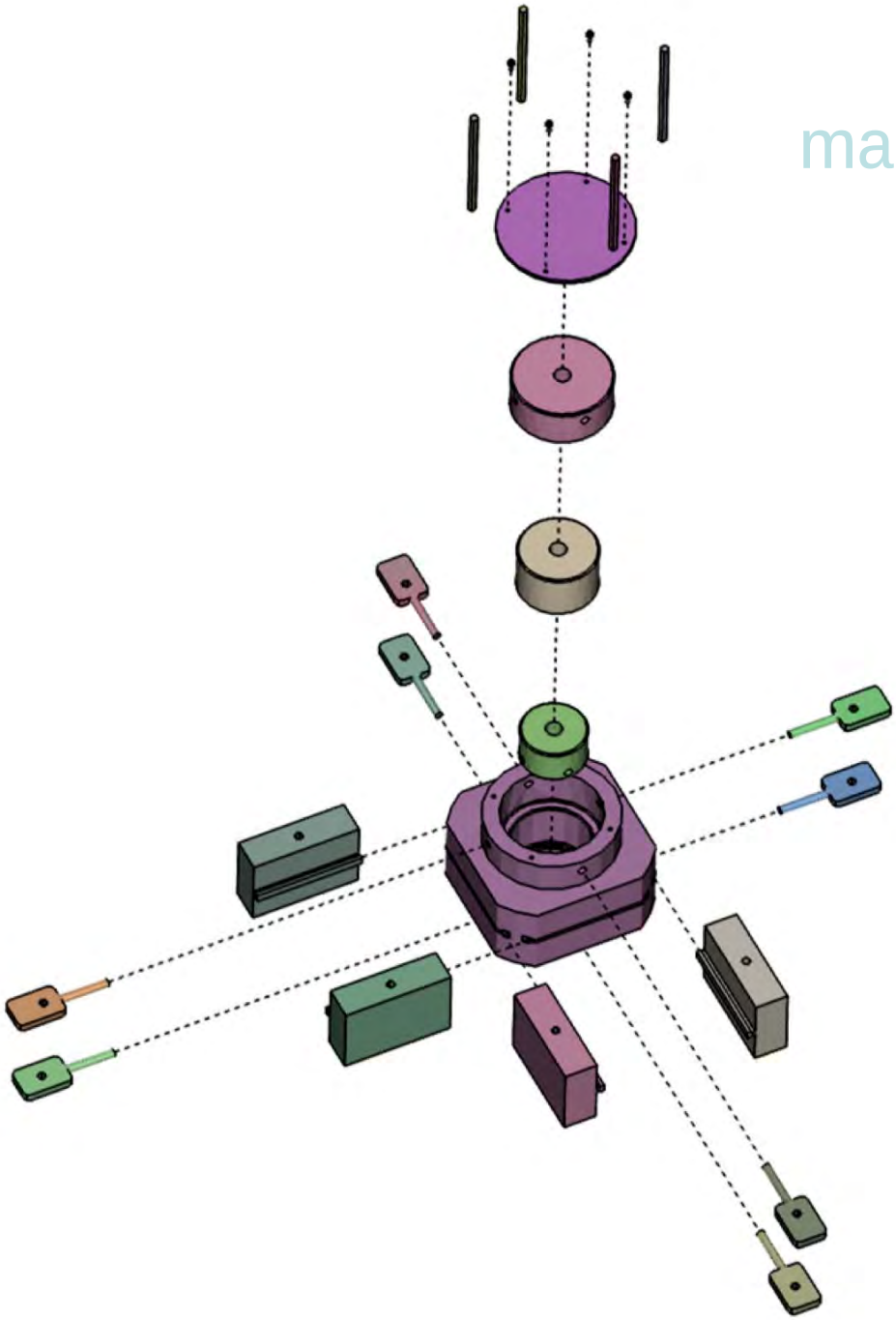

Examiner Viewer
0
1
/491
NMDA.C1B
fast_sp.U
NMDA.C2
fast_sp.B
NMDA.C2B
Ca
Glu
ALL
MESHES
VOLUME_MOLECULES
SURFACE_MOLECULES
search
TOGGLE SELECT
TOGGLE VISIBLE



presenting your data descriptively…

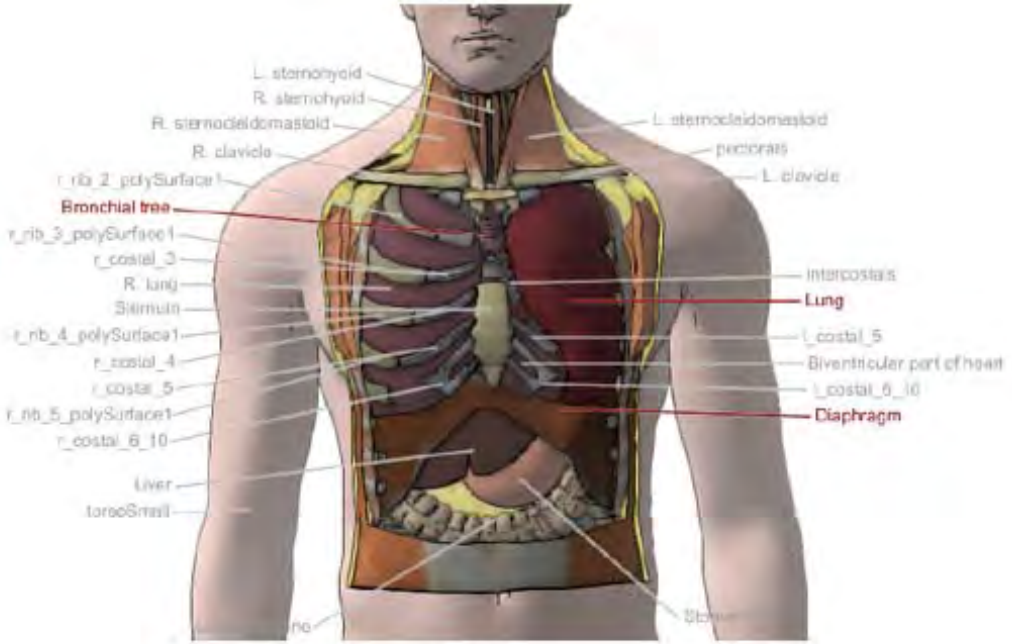
advanced: *techniques*

interactive cutaways

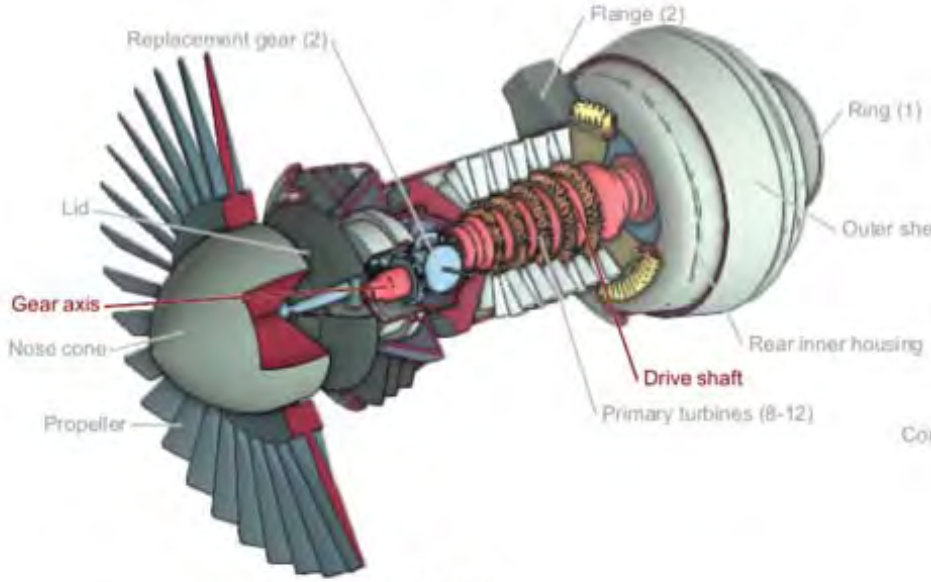


maneesh agrawala

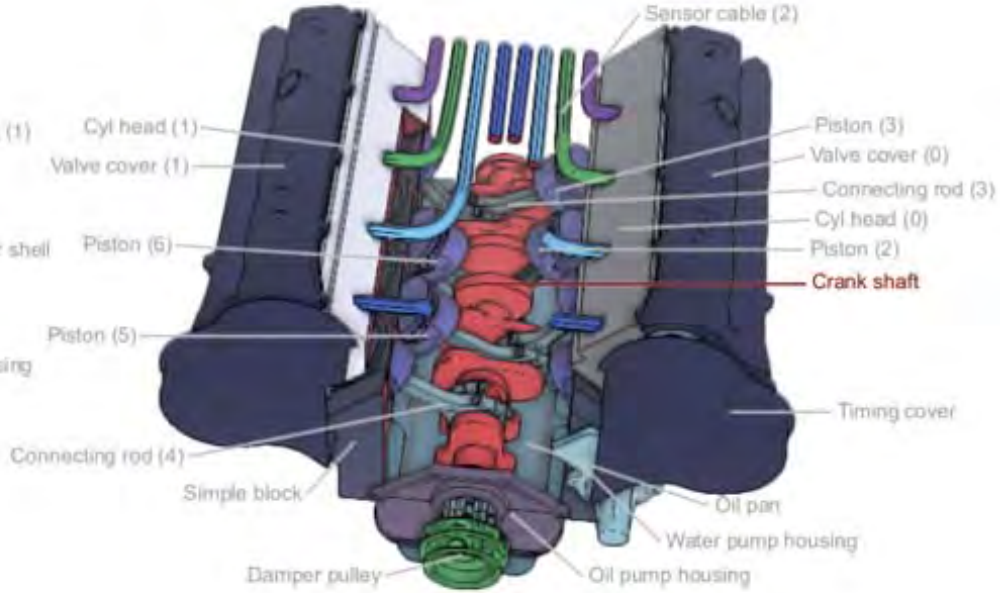
(a) Arm



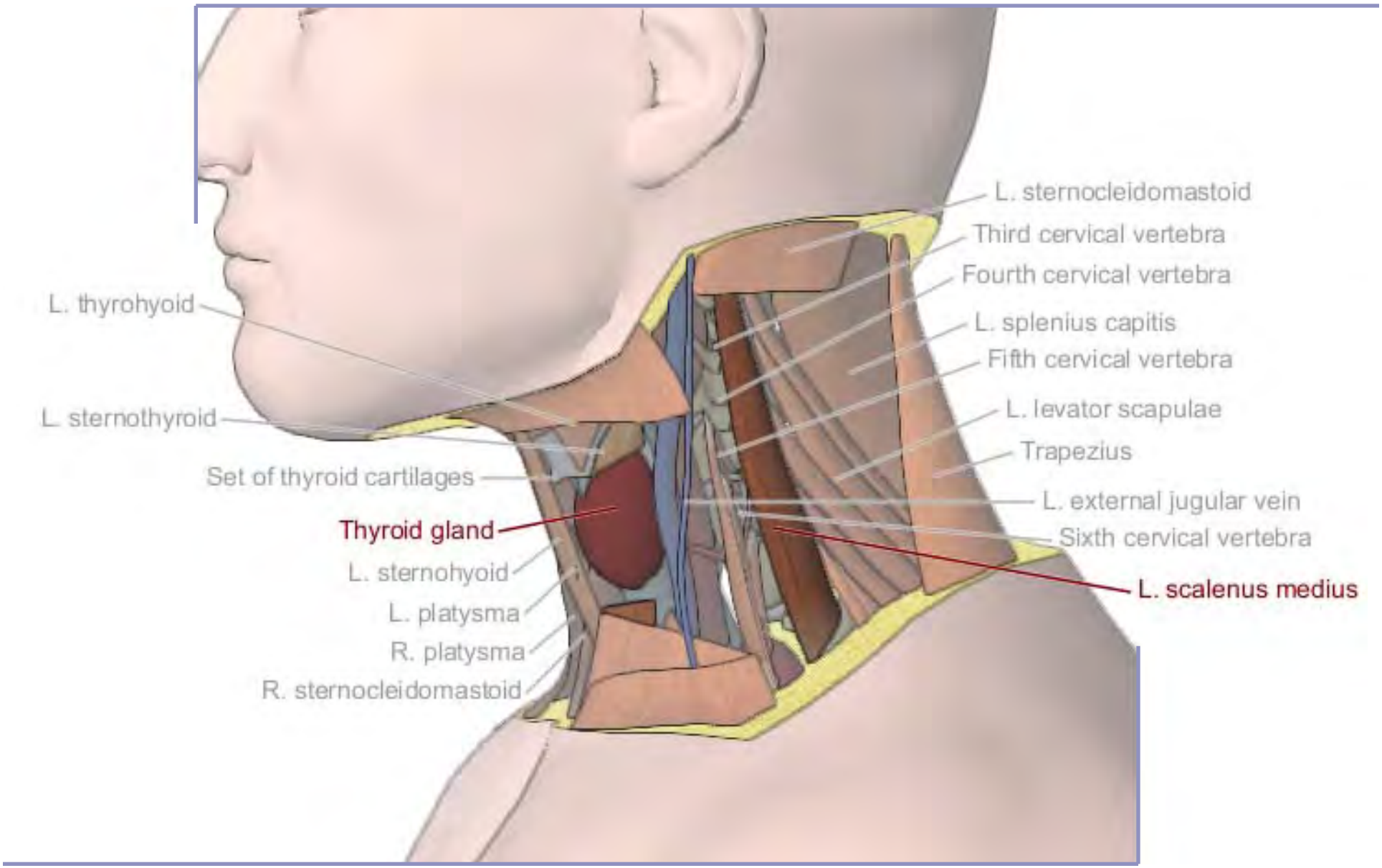
(b) Thorax

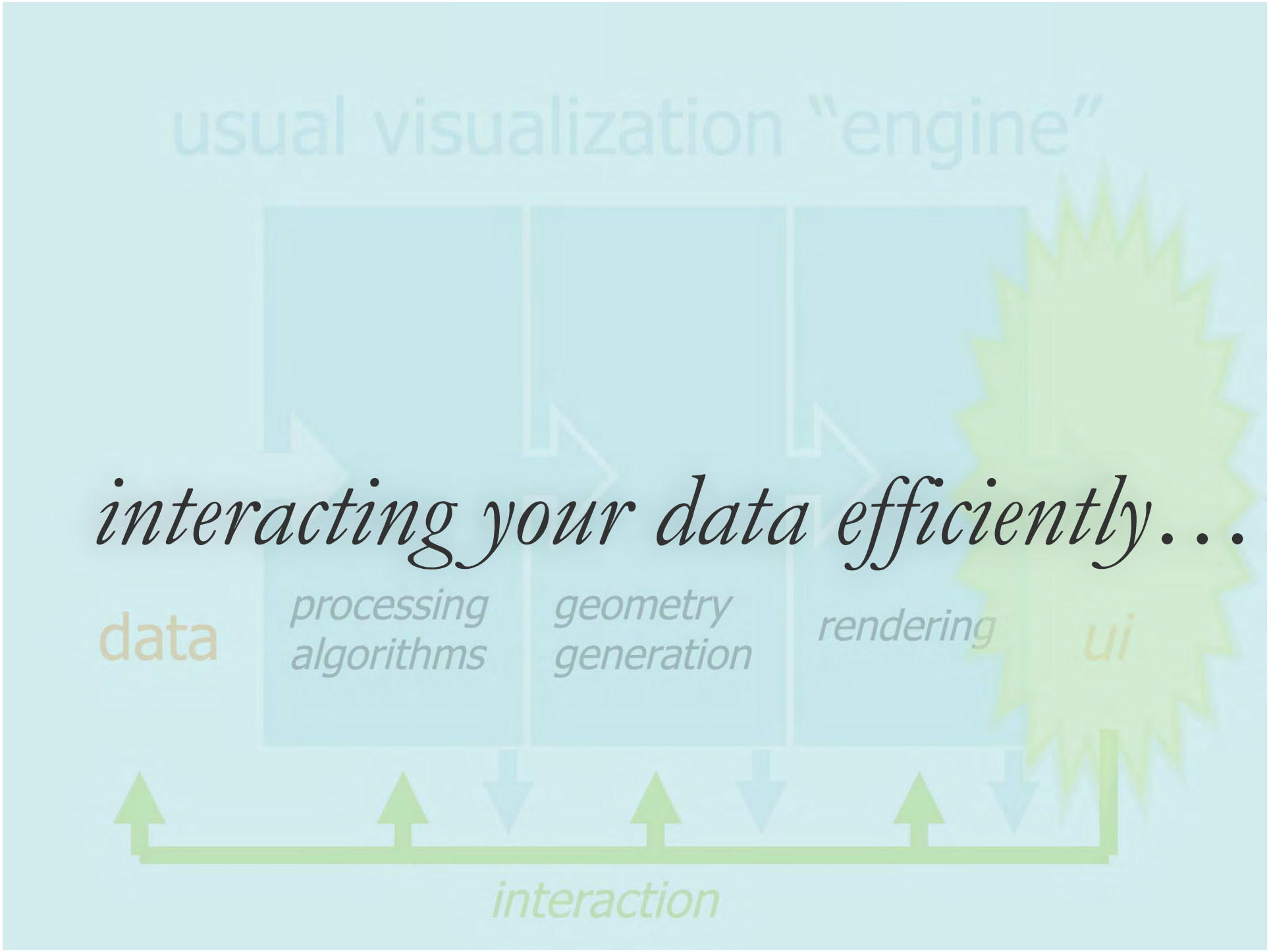


(c) Turbine



(d) Engine


L. thyrohyoid
L. sternothyroid
Set of thyroid cartilages
Thyroid gland
L. sternohyoid
L. platysma
R. platysma
R. sternocleidomastoid
L. sternocleidomastoid
Third cervical vertebra
Fourth cervical vertebra
L. splenius capitis
Fifth cervical vertebra
L. levator scapulae
Trapezius
L. external jugular vein
Sixth cervical vertebra
L. scalenus medius

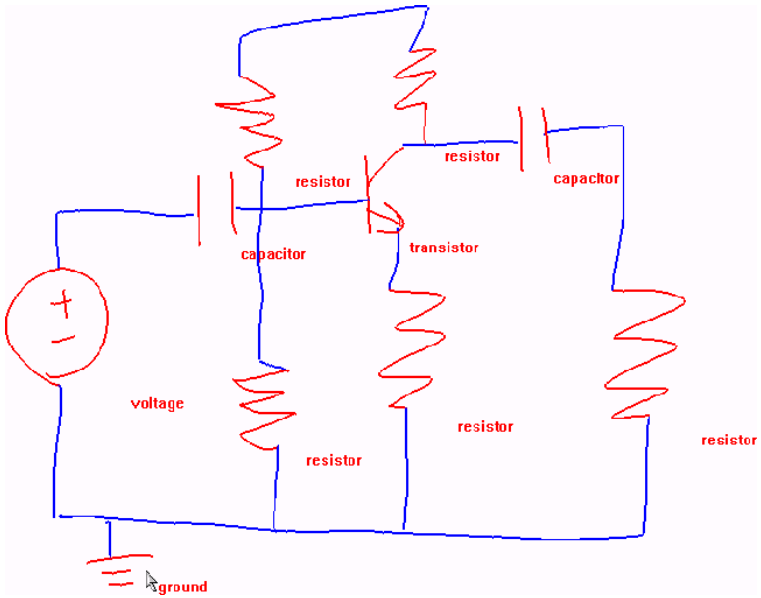


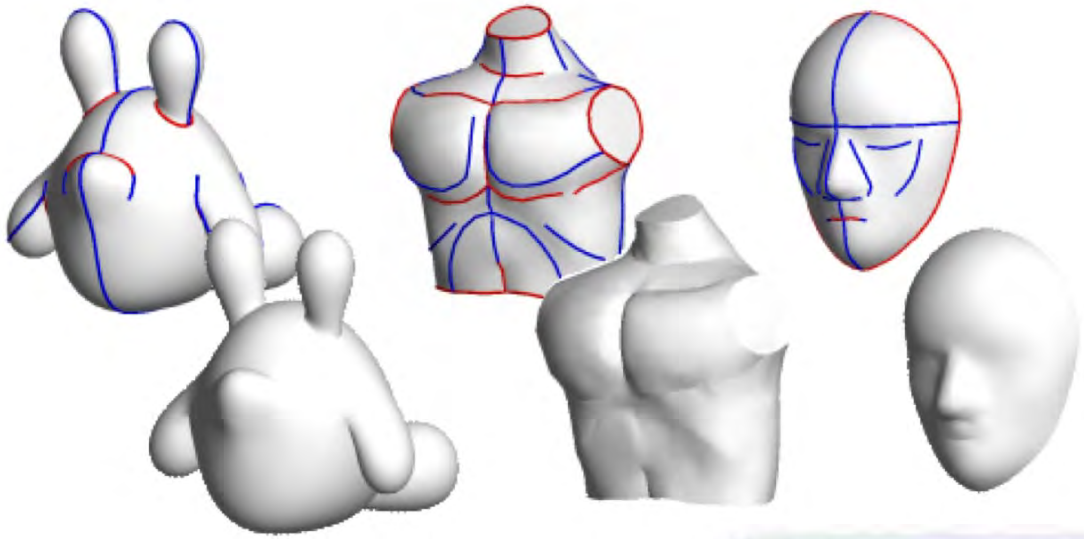
interacting your data efficiently…

advanced: *techniques*
sketch-based uis

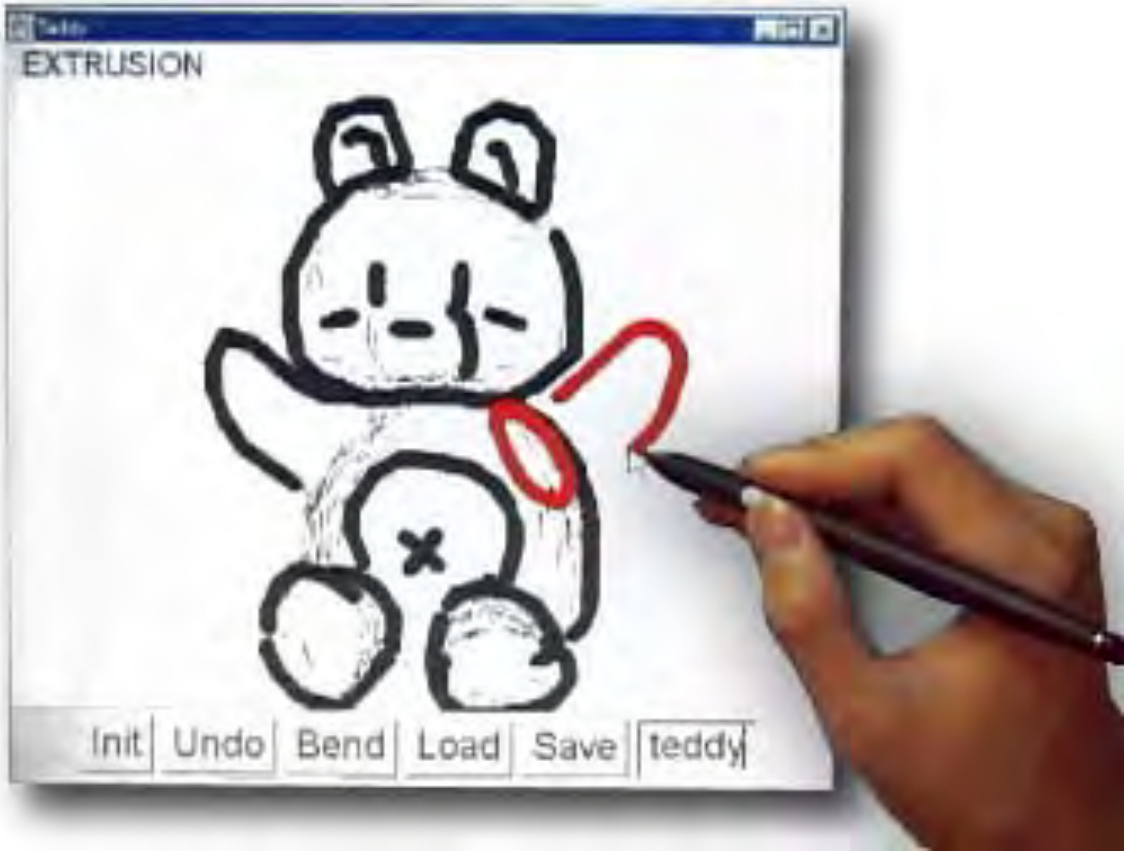
sketch based interfaces

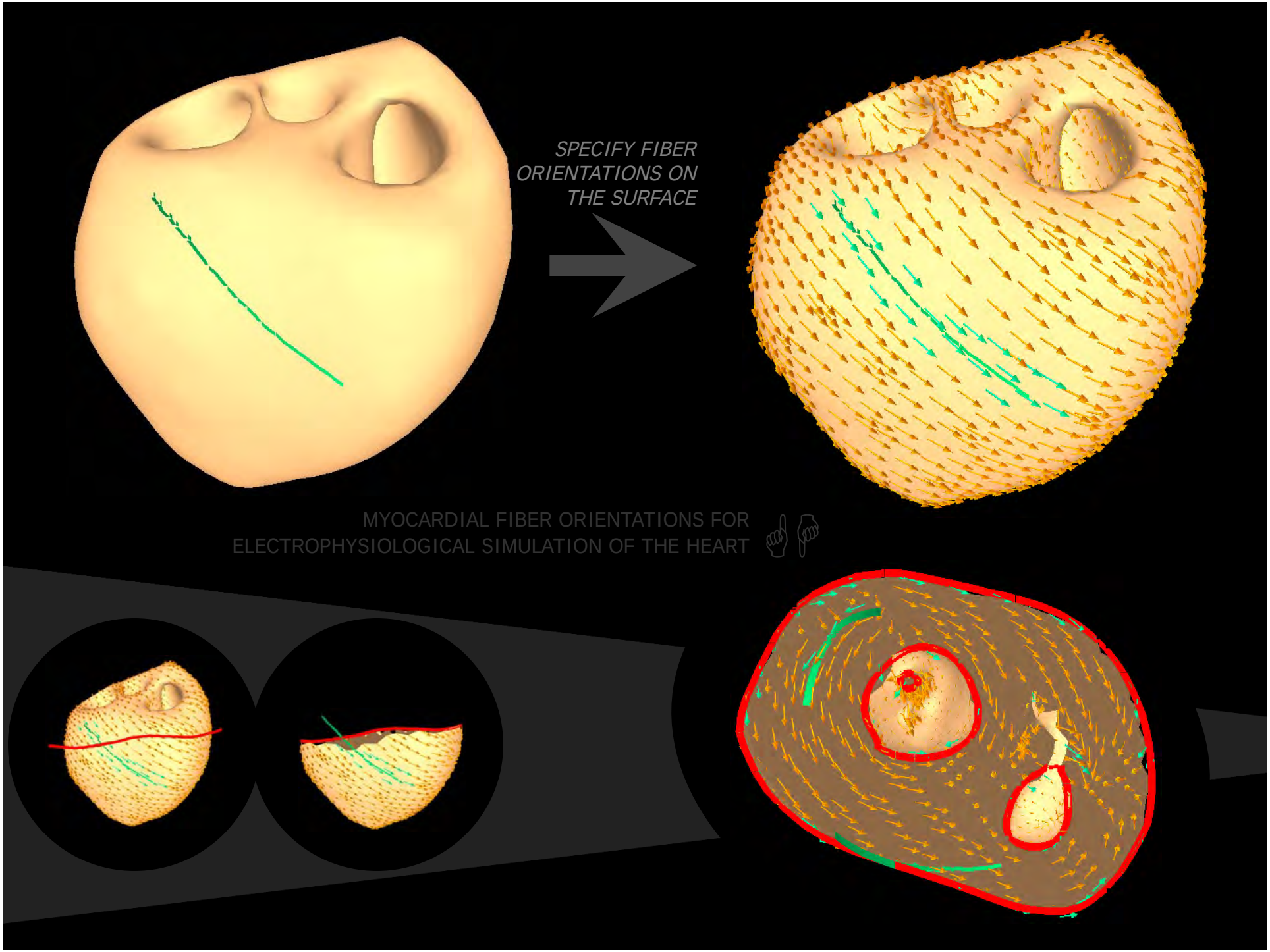
- buttons/menus -> sketch lines
  - i.e. direct interaction!
- more intuitive
- more efficient
- more elegant





takeo igarashi




SPECIFY FIBER
ORIENTATIONS ON
THE SURFACE
MYOCARDIAL FIBER ORIENTATIONS FOR
ELECTROPHYSIOLOGICAL SIMULATION OF THE HEART

√ carefully crafted

visual exploration ☛ answer questions

(even if questions are not clearly defined?)

☛ tools address specific task/question

(visualization is bound by tools... right?)

\+ cross referencing

\+ interactivity

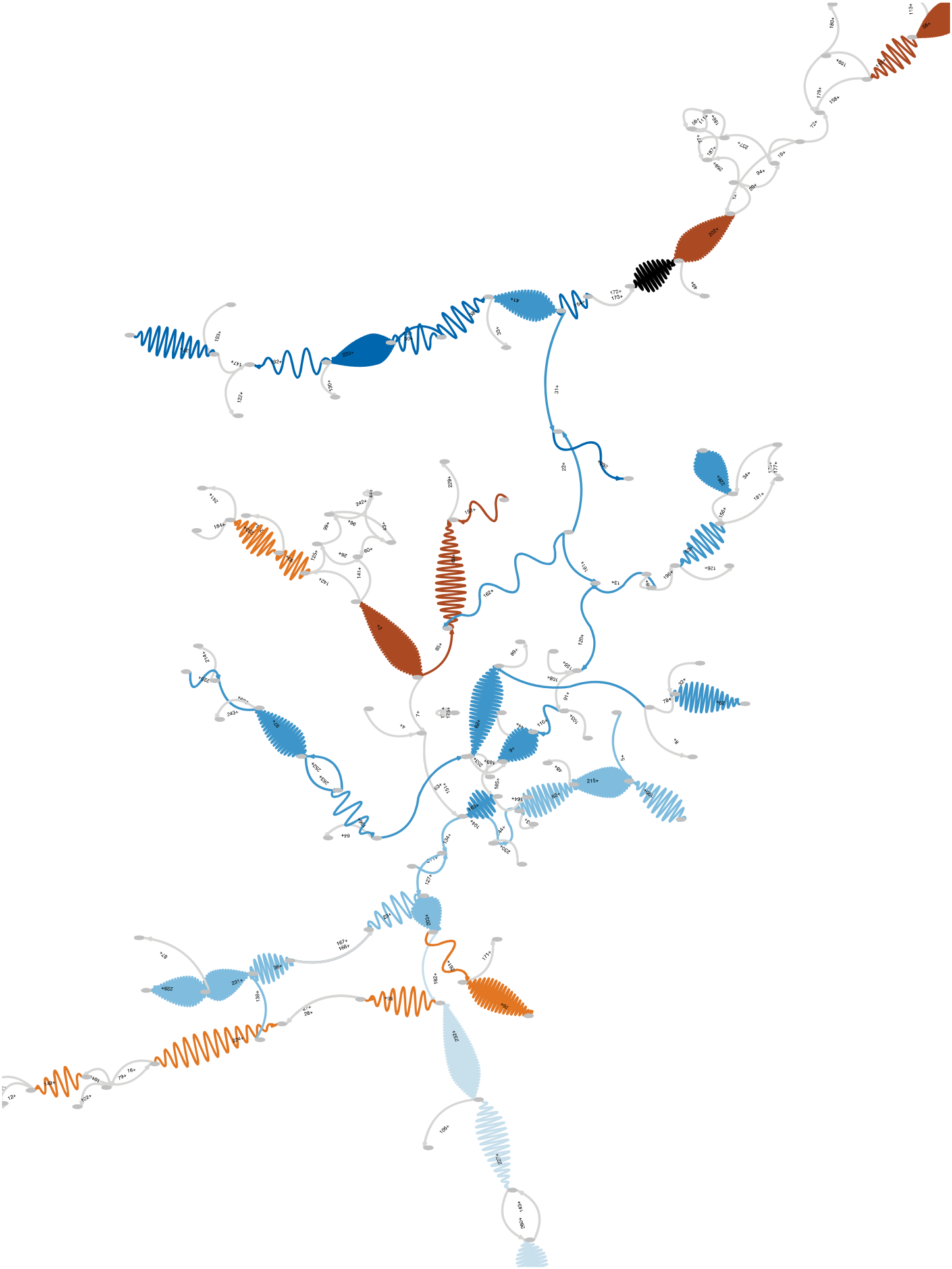
☛ *broader* set of questions addressable

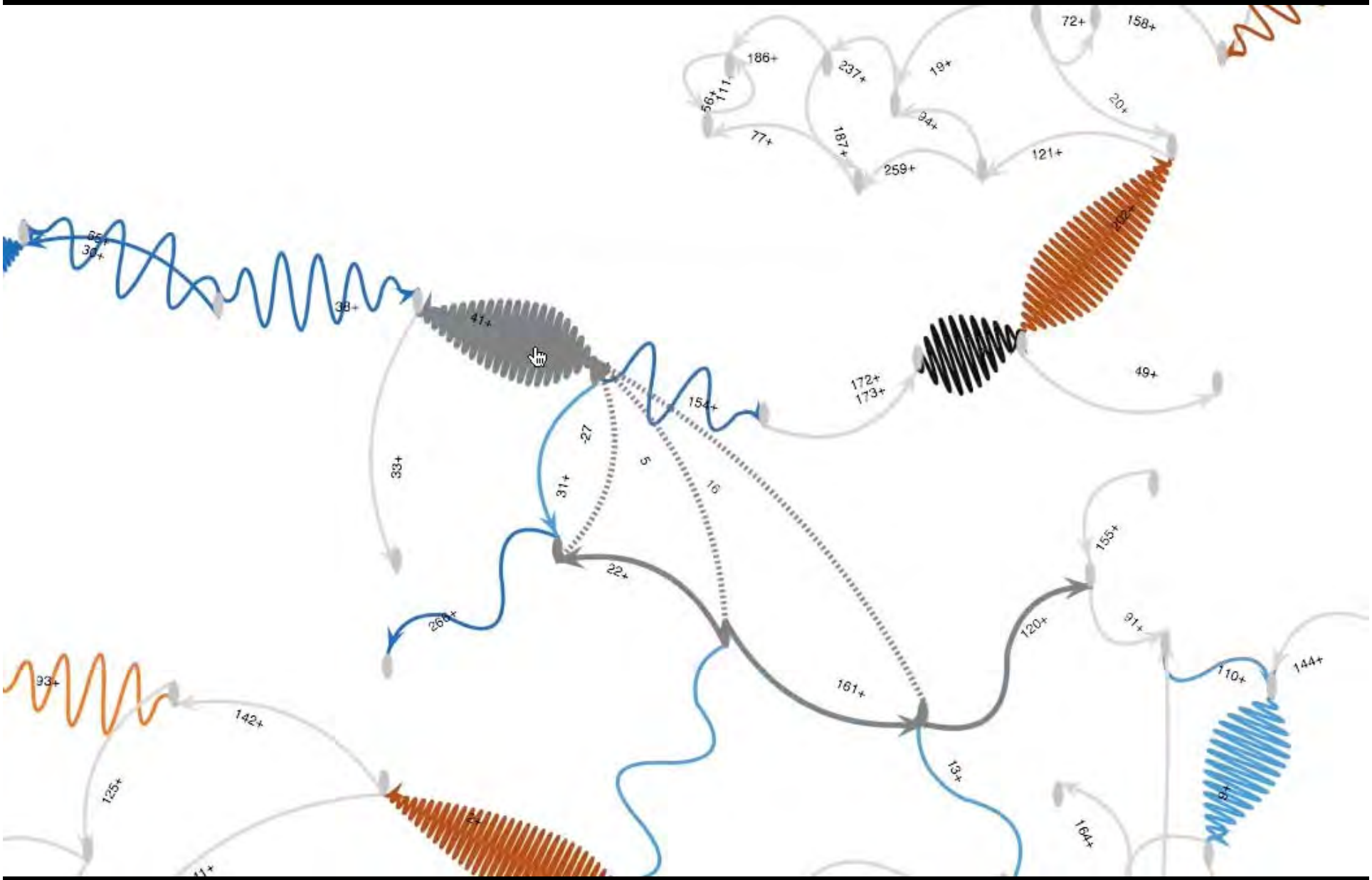
ay/bi199: methods of computational science

visualization

visualization system+techniques

santiago v lombeyda | center for advanced computing research | caltech




72+
158+
186+
237+
19+
56+
111+
20+
34+
77+
187+
259+
121+
202+
38+
41+
172+
173+
49+
154+
27
5
16
33+
31+
22+
155+
266+
120+
91+
93+
142+
161+
110+
144+
125+
13+
164+
24
8+