

# IV121 Vybrané aplikace informatiky v biologii

## 3D počítačová grafika

Katedra informačních technologií  
Masarykova Univerzita Brno

Jaro 2012

Tento projekt je spolufinancován Evropským sociálním fondem a státním rozpočtem České republiky.



### Stringologie

Úvod

Základní pojmy

Základní algoritmy

Algoritmus využívající analýzu  
hledaného motivu

Algoritmus využívající analýzu  
prohledávaného řetězce

### Hledání opakování

Tandemové opakování

Palindromy

### Srovnávání dvou sekvencí

DP - Needleman-Wunsch

Vylepšení pro maximálně k chyb

Burrows-Wheeler transform

# 3D Počítačová Grafika (nebo Geometrie)

- ***modelování scén***

  - SDL (scene description language)

- ***vizualizace scén (rendering)***

  - rasterizace

  - „raytracing“

- ***zajímavé koncepty***

  - CSG (constructive solid geometry)

  - skriptování scén

  - příklad generování realistických stromů a keřů

# SDL – Scene Description Languages

***VRML/X3D***

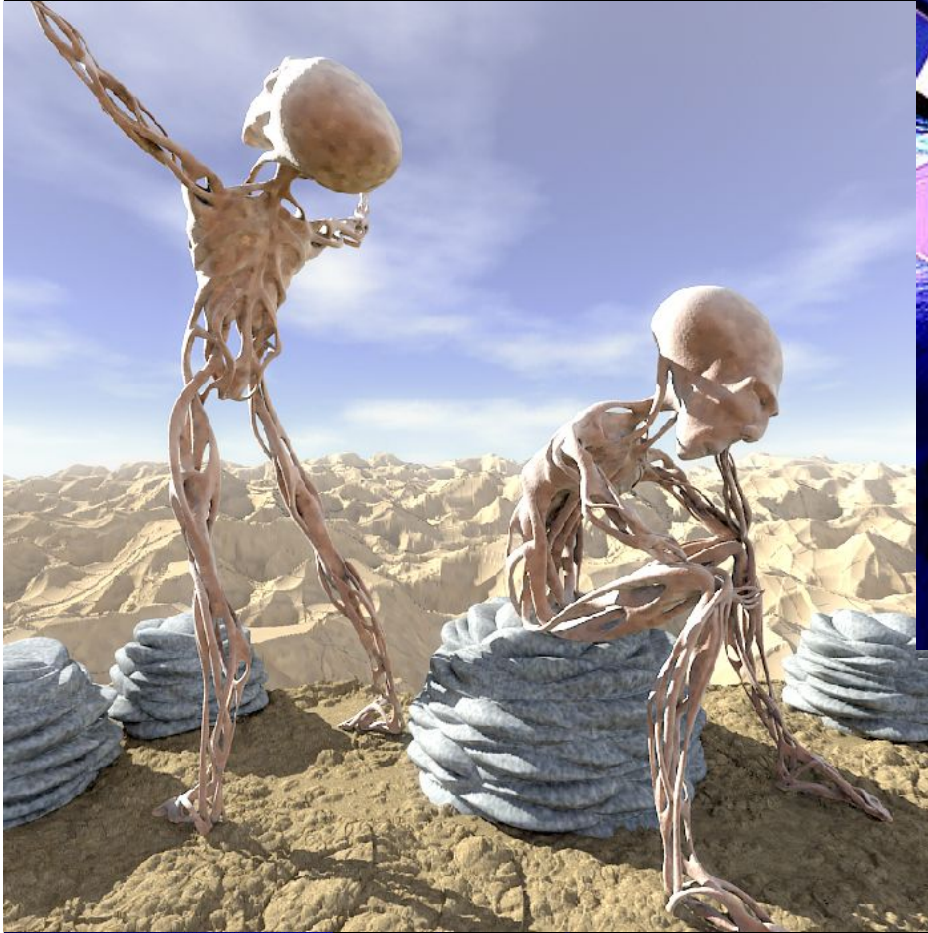
***3DMLW***

***POV-Ray SDL***

***Renderman shading language***

[http://en.wikipedia.org/wiki/Scene\\_description\\_language](http://en.wikipedia.org/wiki/Scene_description_language)

# Co je RenderMan?



Entropy image contest winner by Claude Schitter



MonstersInc by Pixar



A Bug's Life by Pixar



# Co je RenderMan?

Autorem je společnost Pixar (1987)

Něco jako PostScript pro 3D

- Scene Description Language

není modelovacím programem

není renderingovým programem

Rozhraním mezi modelováním a renderingem

# Příklad Bytestream kódu pro RenderMan Interface

```
Display "RenderMan" "framebuffer"  
  "rgb"  
Format 256 192 1  
WorldBegin  
Surface "constant"  
Polygon "P" [0.5 0.5 0.5 0.5 -0.5  
  0.5 -0.5 -0.5 0.5 -0.5 0.5 0.5]  
WorldEnd
```

# API

```
#include <ri.h>

RtPoint Square[4] = { {.5, .5, .5}, {.5, -.5, .5}, {-.5, -.5, .5},
                      {-.5, .5, .5} };

main(void) {
    RiBegin(RI_NULL);    /* Start the renderer */
    RiDisplay("RenderMan", RI_FRAMEBUFFER, "rgb", RI_NULL);
    RiFormat((RtInt) 256, (RtInt) 192, 1.0);
    RiWorldBegin();

        RiSurface("constant", RI_NULL);
        RiPolygon( (RtInt) 4,          /* Declare the square */
                  RI_P, (RtPointer) Square, RI_NULL);
    RiWorldEnd();

    RiEnd();            /* Clean up */
}
```

# RenderMan Shading Language

```
surface clouds(float vfreq = .8 )
{
    float sum ;
    float i;
    color white = color(1.0, 1.0, 1.0);
    point Psh = transform("shader", P);

    sum = 0;
    freq = vfreq;
    for (i = 0; i < 6; i = i + 1) {
        sum = sum + 1/freq * abs(.5 - noise(freq * Psh));
        freq = 2 * freq;
    }
    Ci = mix(Cs, white, sum*4.0);
    Oi = 1.0;          /* Always make the surface opaque */
}
```



# RIB s použitím "Shader" kódu

```
Display "RenderMan" "framebuffer"  
"rgb"
```

```
Format 256 192 1
```

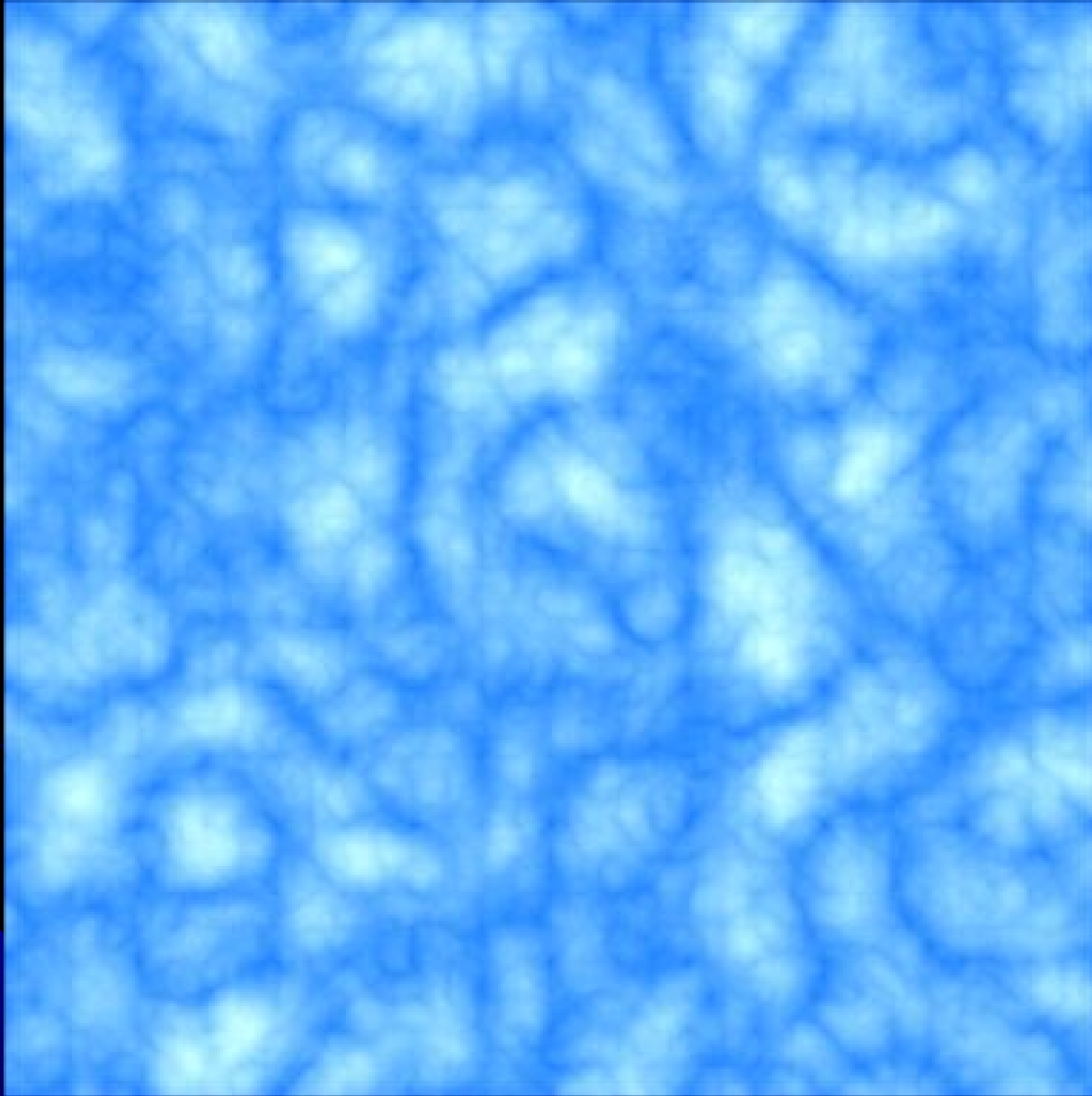
```
WorldBegin
```

```
Surface "clouds"
```

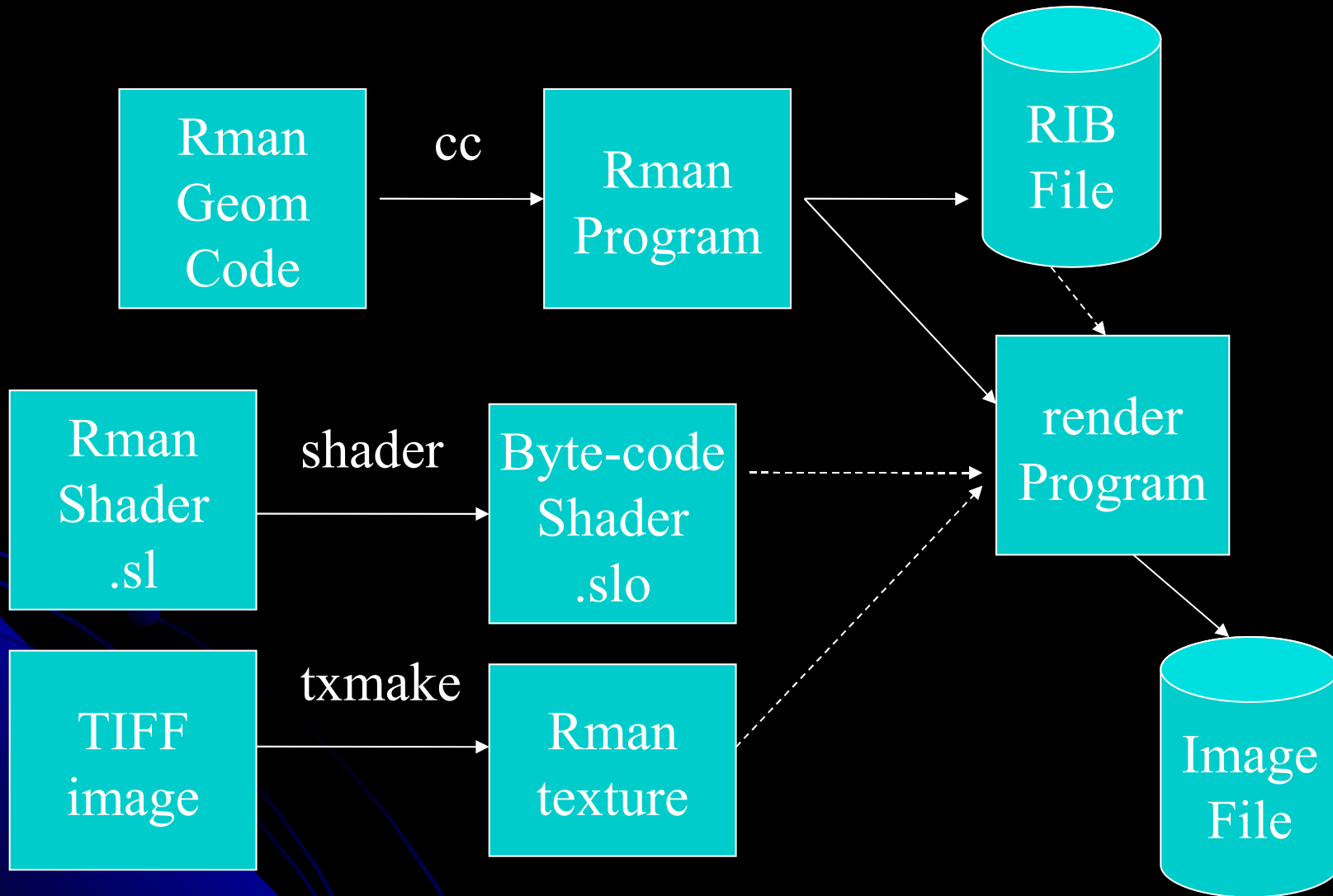
```
Polygon "P" [0.5 0.5 0.5 0.5 -0.5  
0.5 -0.5 -0.5 0.5 -0.5 0.5 0.5]
```

```
WorldEnd
```

# Result



# Součásti systému RenderMan



# „Rendering“

## - *rasterizace*

Vlastnosti všech bodů v prostoru/modelu jsou lokálně definovány

## - *“raytracing“*

Vlastnosti bodů jsou ovlivněny globálně všemi ostatními body ve scéně/modelu.

Dokáže správně vykreslit transparenční, refrakční světla a podobné efekty.



# Rendering Pipeline - OpenGL

Aaron Bloomfield

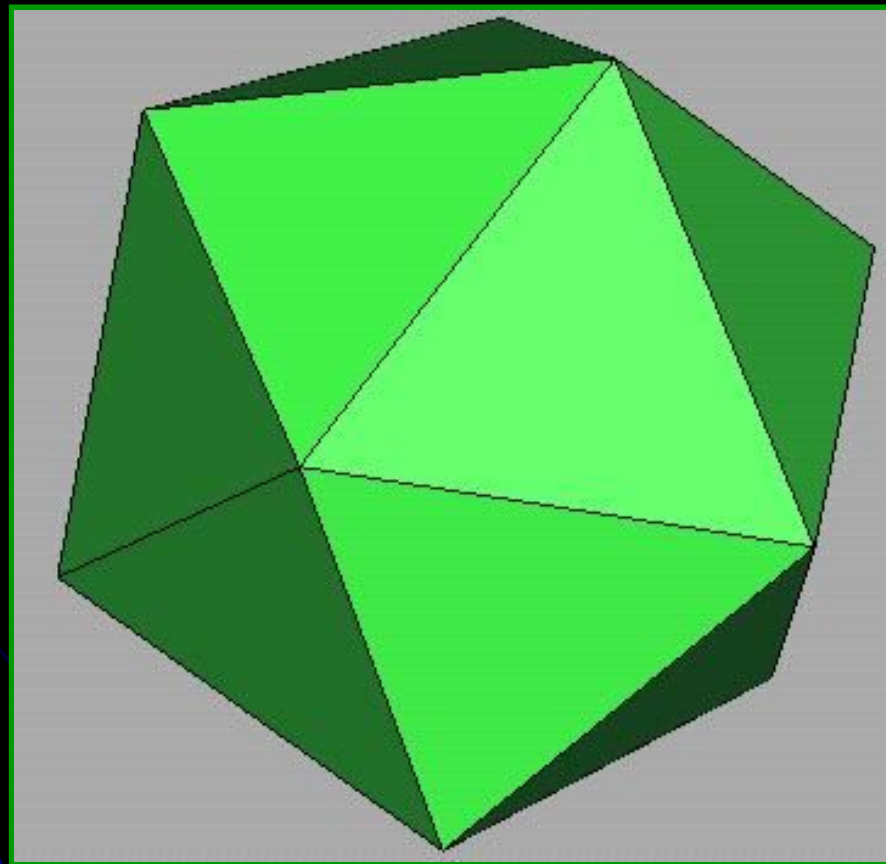
CS 445: Introduction to Graphics

Fall 2006

(Slide set originally by Greg Humphreys)

# 3D Polygon Rendering

Many applications use rendering of 3D polygons with direct illumination



# 3D Polygon Rendering

Many applications use rendering of 3D polygons with direct illumination



# 3D Rendering Pipeline

## 3D Geometric Primitives

Modeling  
Transformation

Lighting

Viewing  
Transformation

Projection  
Transformation

Clipping

Scan  
Conversion

Image

This is a pipelined sequence of operations to draw a 3D primitive into a 2D image

(this pipeline applies only for direct illumination)



# Viewing Transformation

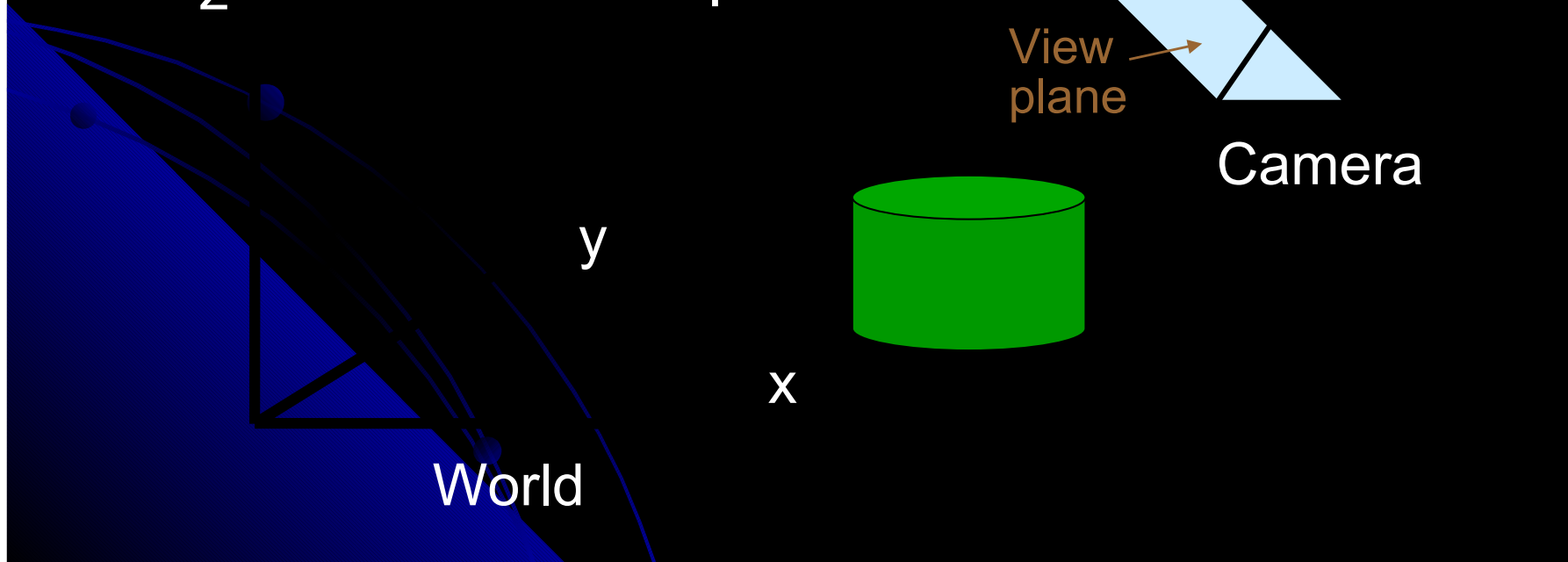
Mapping from world to camera coordinates

Eye position maps to origin

Right vector maps to X axis

Up vector maps to Y axis

$z$  Back vector maps to Z axis



# Projection

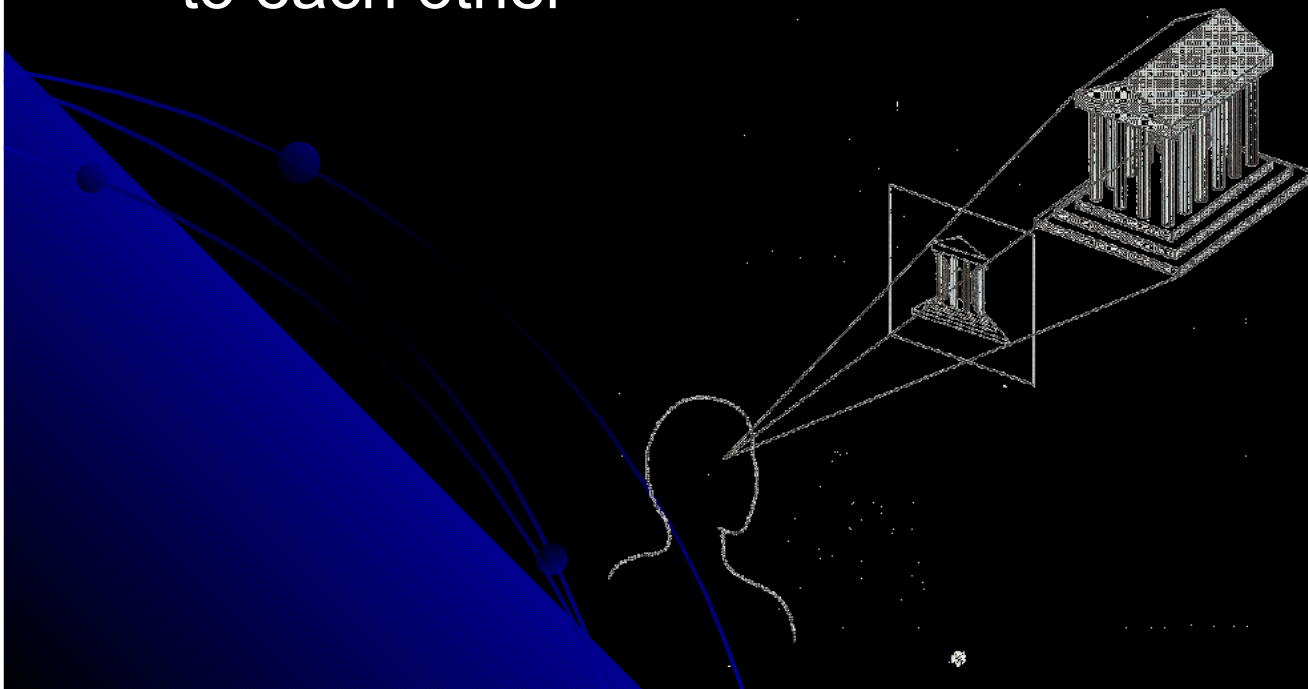
General definition:

Transform points in  $n$ -space to  $m$ -space ( $m < n$ )

In computer graphics:

Map 3D camera coordinates to 2D screen coordinates

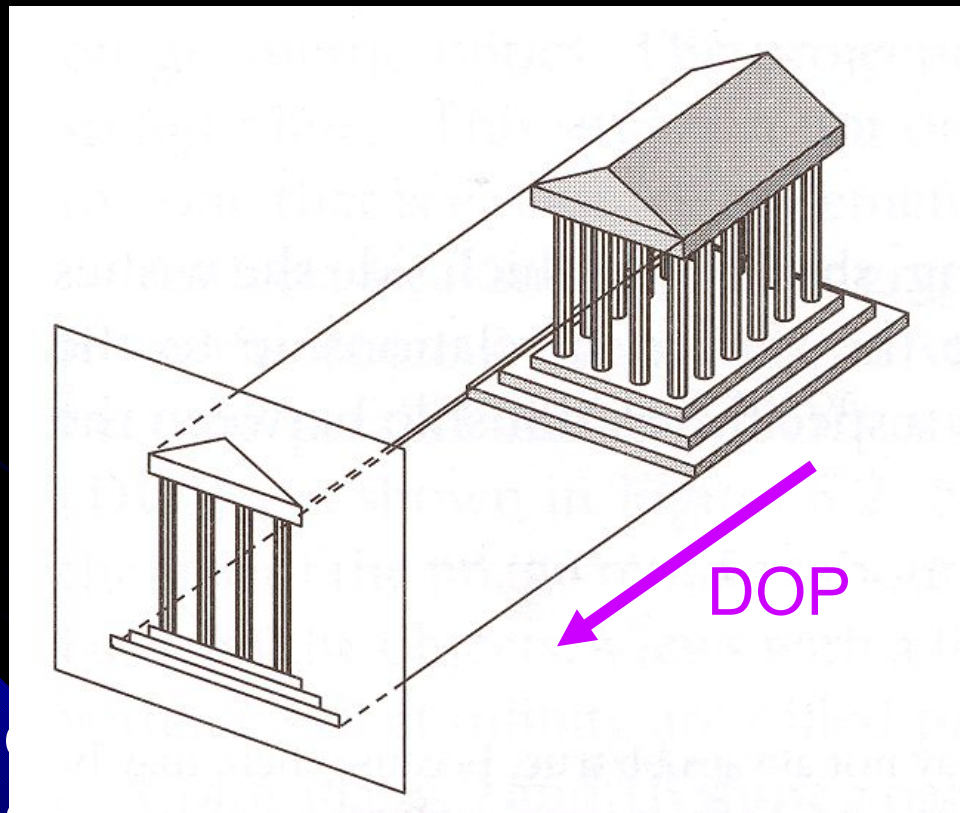
For perspective transformations, no two “rays” are parallel to each other



# Parallel Projection

Center of projection is at infinity

Direction of projection (DOP) same for all points

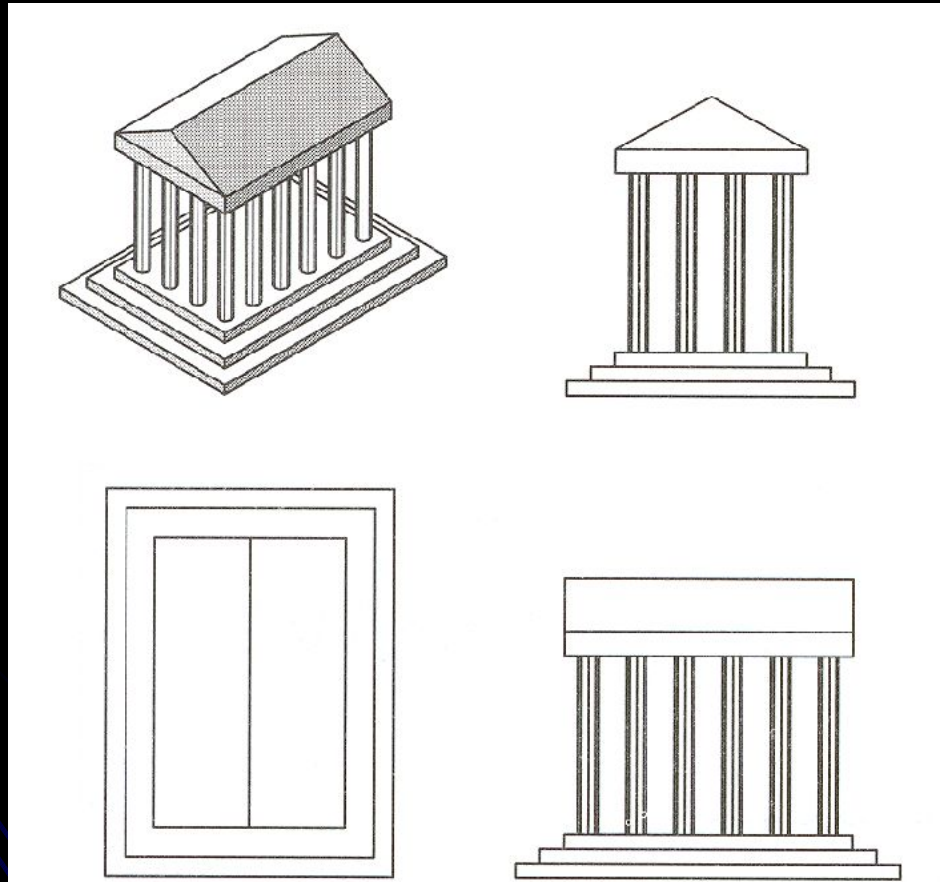


View  
Plane

Angel Figure 5.4<sub>35</sub>

# Orthographic Projections

DOP perpendicular to view plane



Top

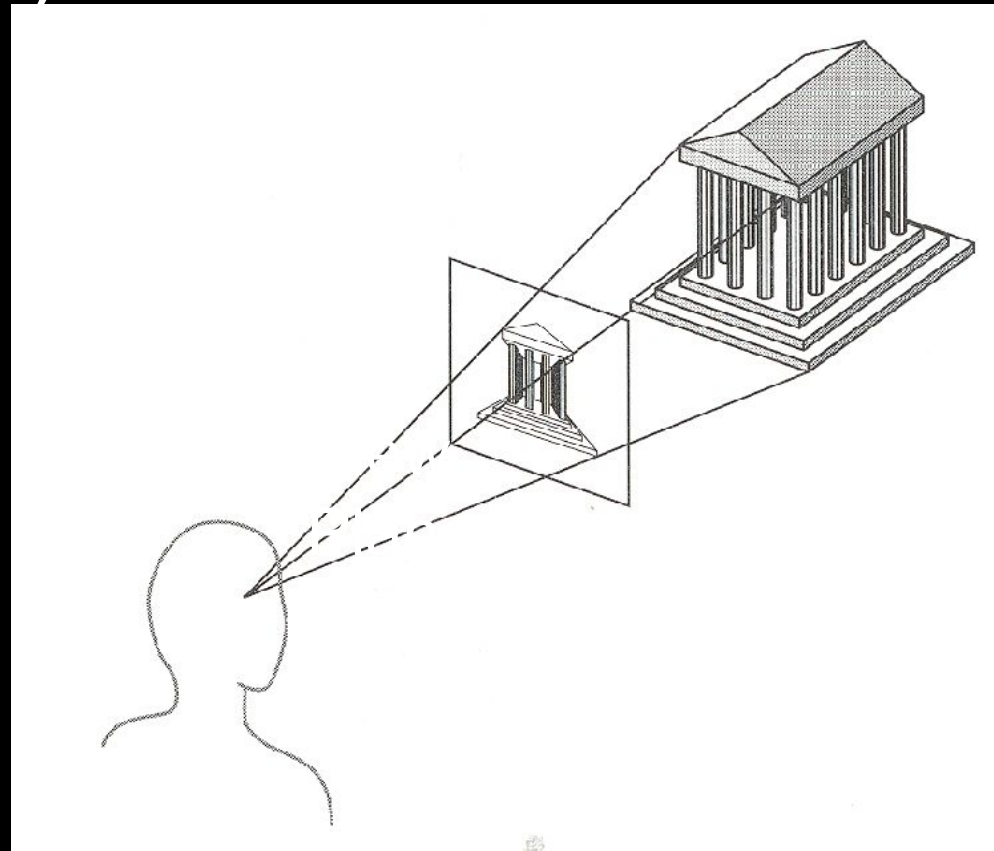
Side



# Perspective Projection

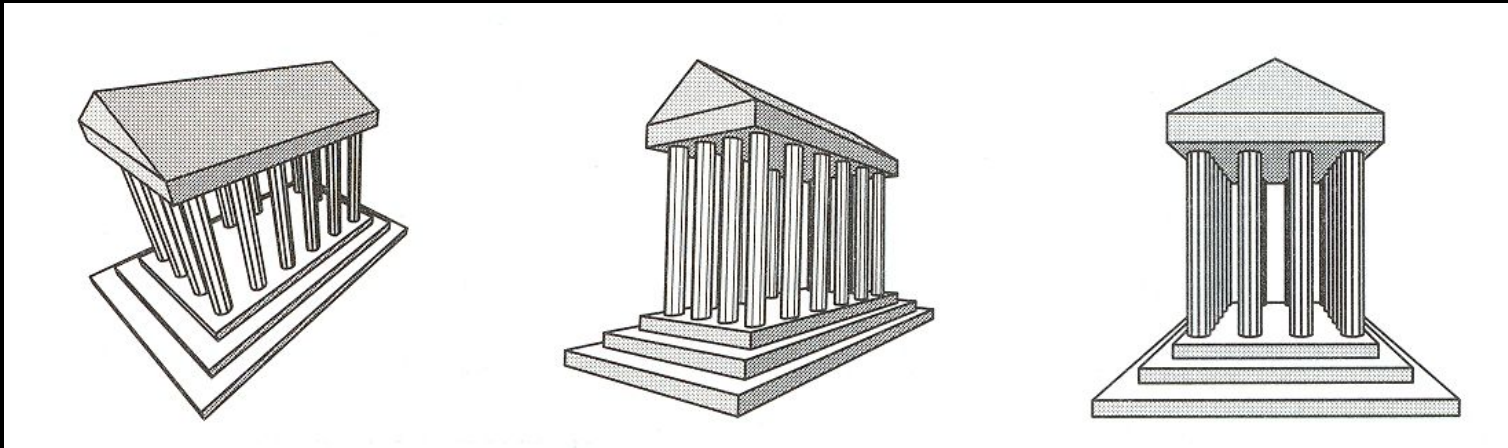
Map points onto “view plane” along  
“projectors” emanating from “center of  
projection” (COP)

Center of  
Projection



# Perspective Projection

How many vanishing points?



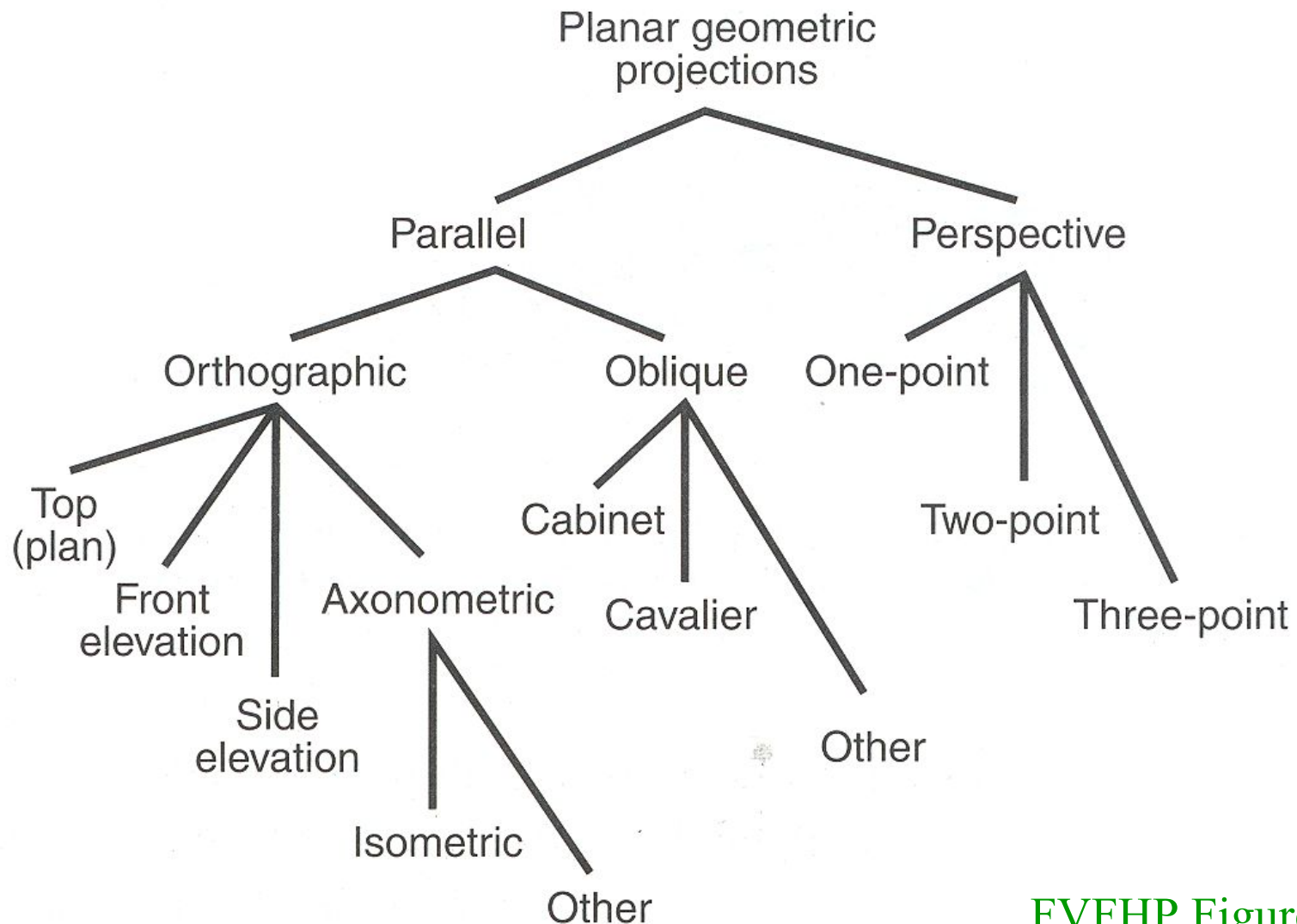
3-Point  
Perspective

2-Point  
Perspective

1-Point  
Perspective

- The difference is how many of the three principle directions are parallel/orthogonal to the projection plane

# Taxonomy of Projections

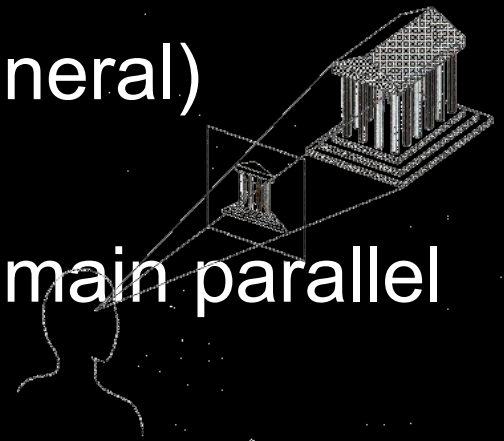


FVFHP Figure 6.10

# Perspective vs. Parallel

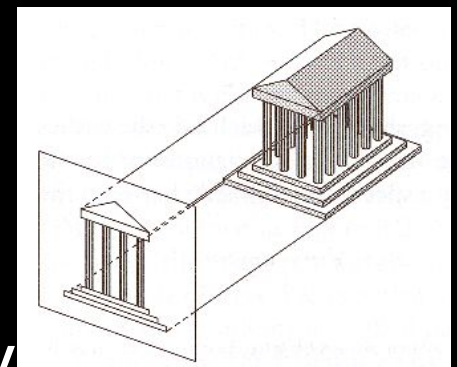
## Perspective projection

- + Size varies inversely with distance - looks realistic
- Distance and angles are not (in general) preserved
- Parallel lines do not (in general) remain parallel



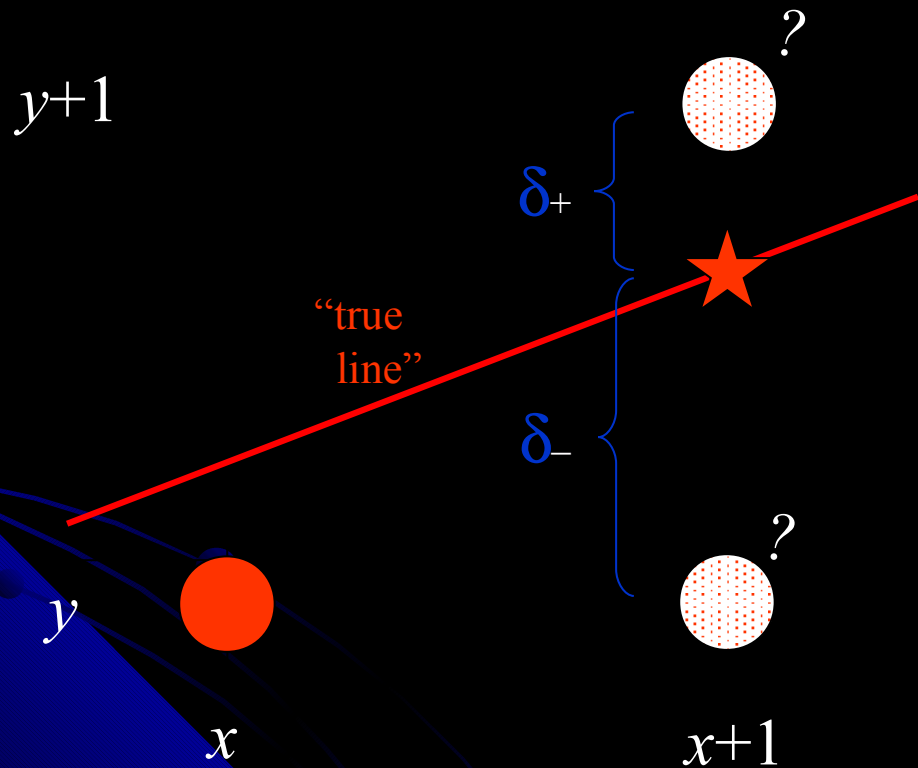
## ■ Parallel projection

- + Good for exact measurements
- + Parallel lines remain parallel
- Angles are not (in general) preserved



# Scan-converting Lines - Toward the Bresenham Algorithm

Special case:  $0 < m < 1, \Delta x > 0$



*imagine pixel centers  
at grid vertices*

*At each step:*

```
x++;  
if (  $\delta_- > \delta_+$  )  
    y++ ;
```

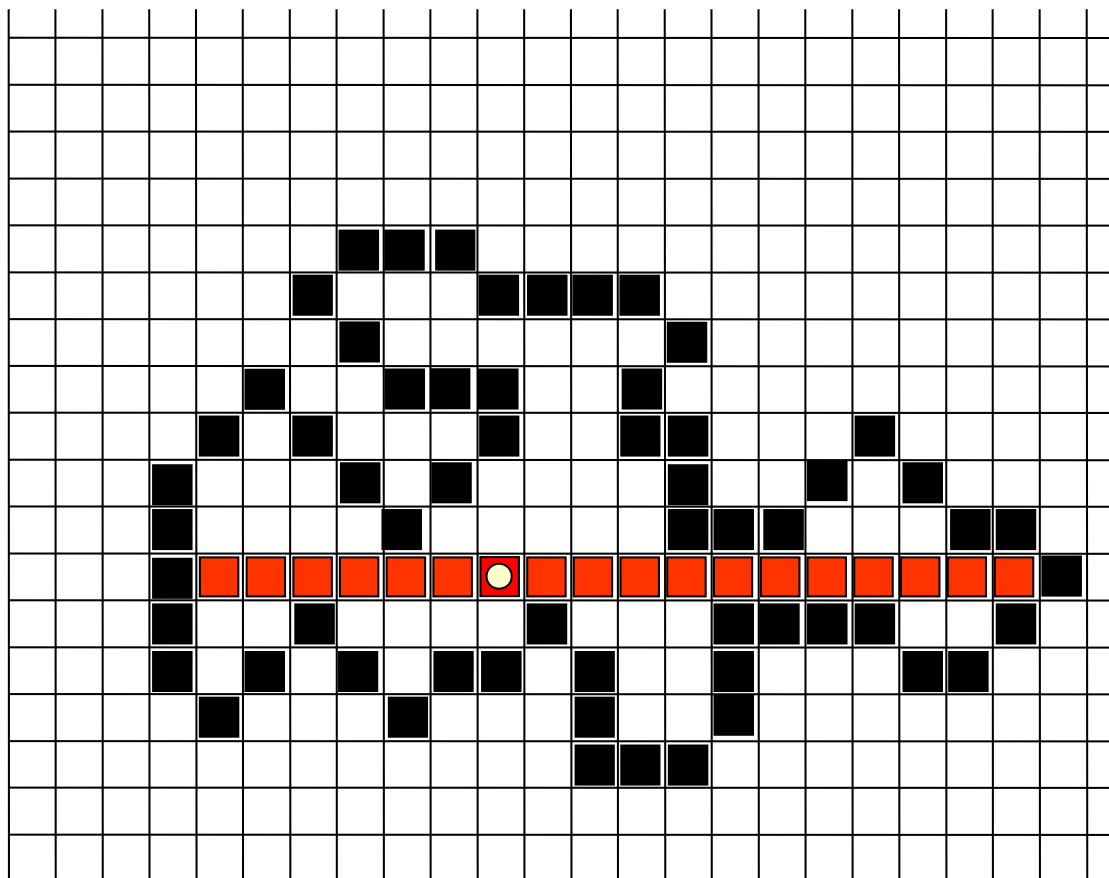
Let  $p = \Delta x (\delta^- - \delta^+)$ . Then:

*At each step:*

```
x++;  
if (  $p > 0$  )  
    y++ ;  
update p ;
```

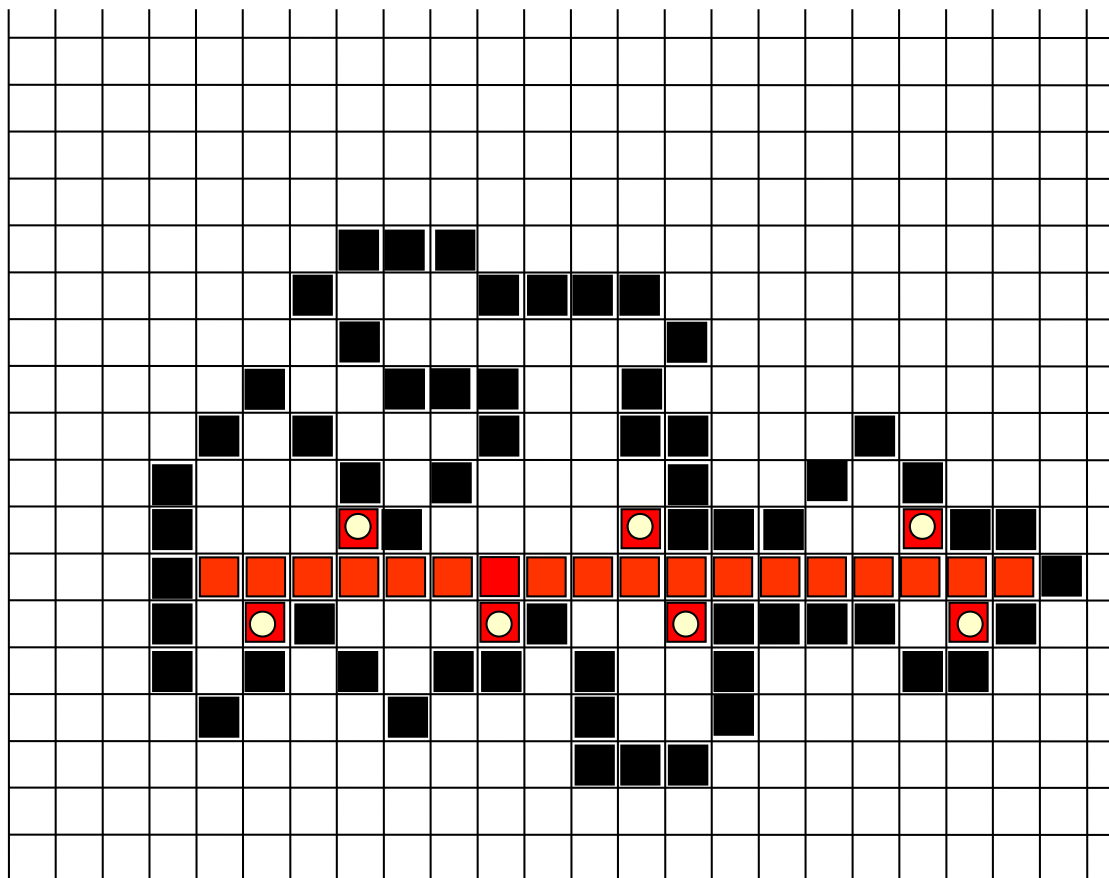
*Can we do this with simple all-int arithmetic? Yes!*





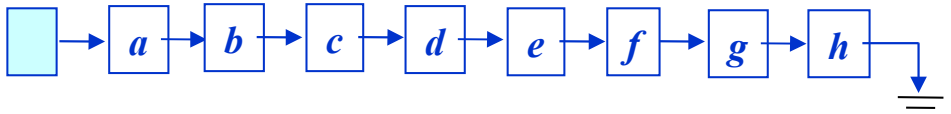
*Fill the  
scan line*

*Push  
seeds for  
the next  
step*

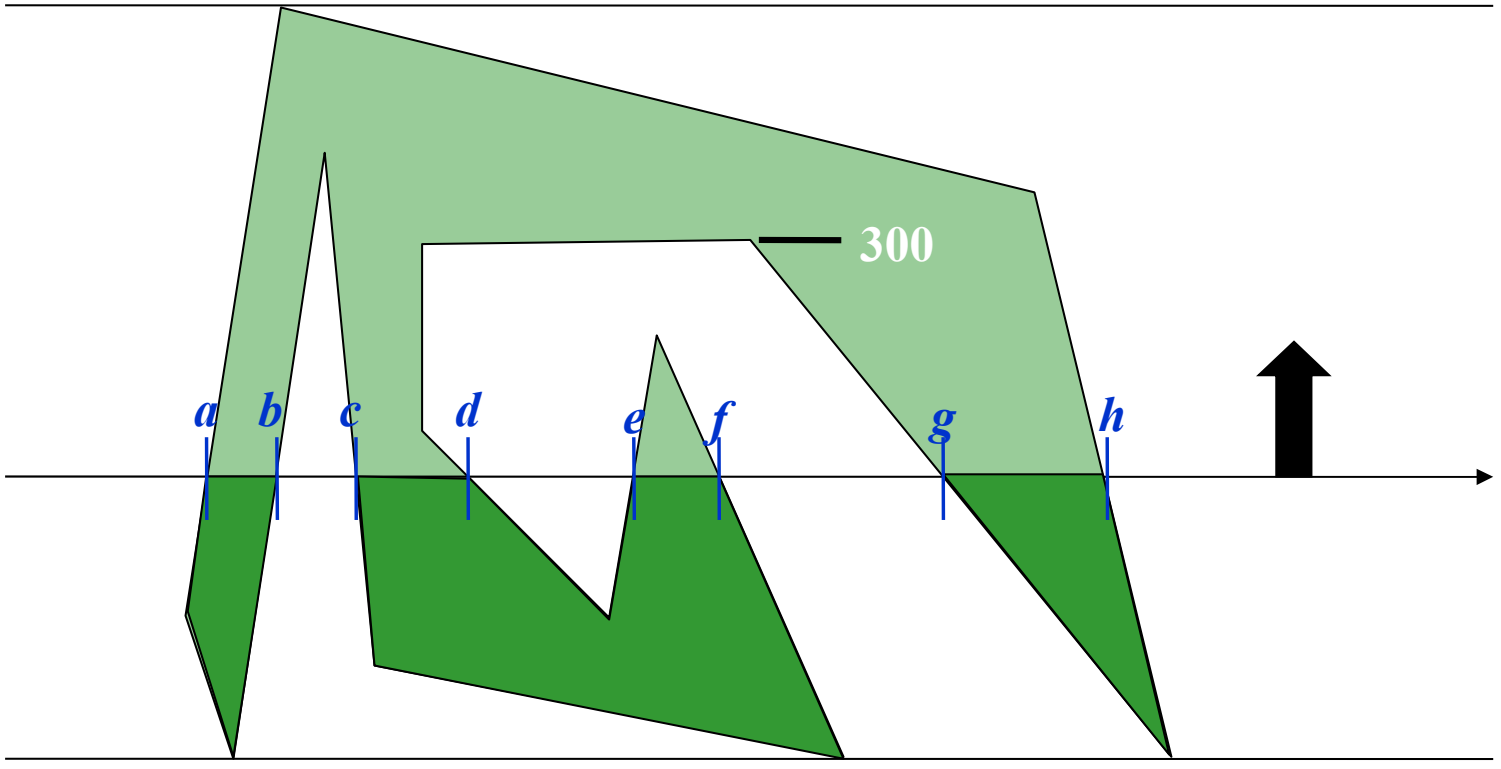


*rightmost left pixels, above & below*





*y*max



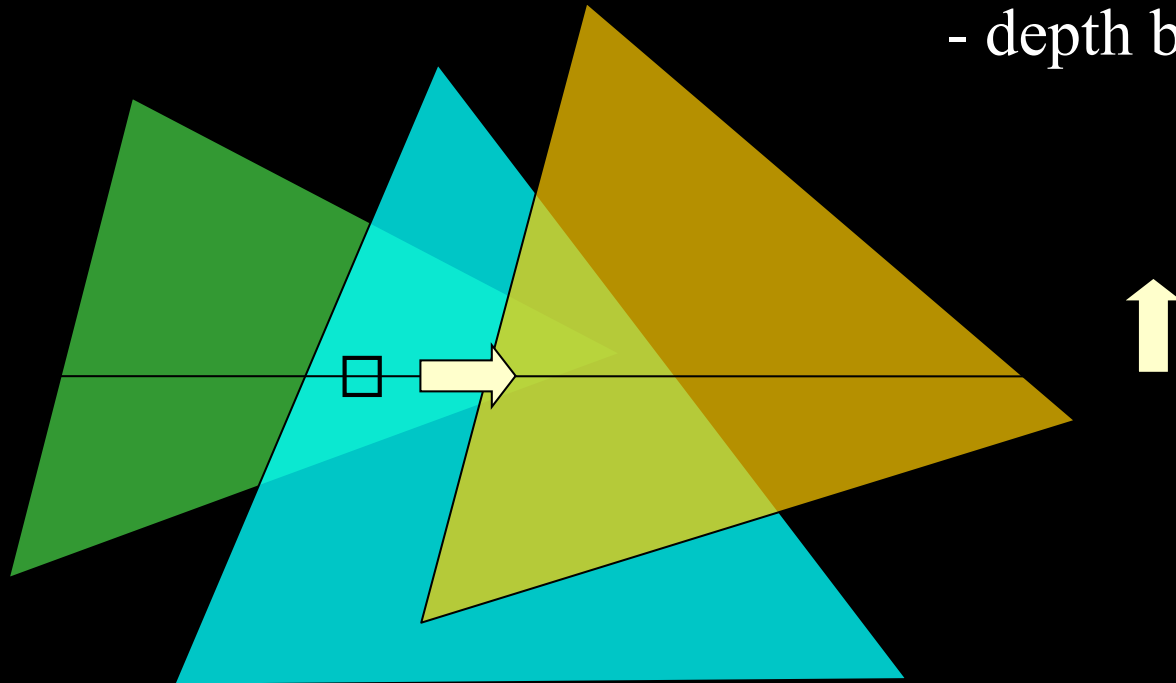
*y*min



# Z-Buffering

## Two buffers

- screen buffer (color)
- depth buffer

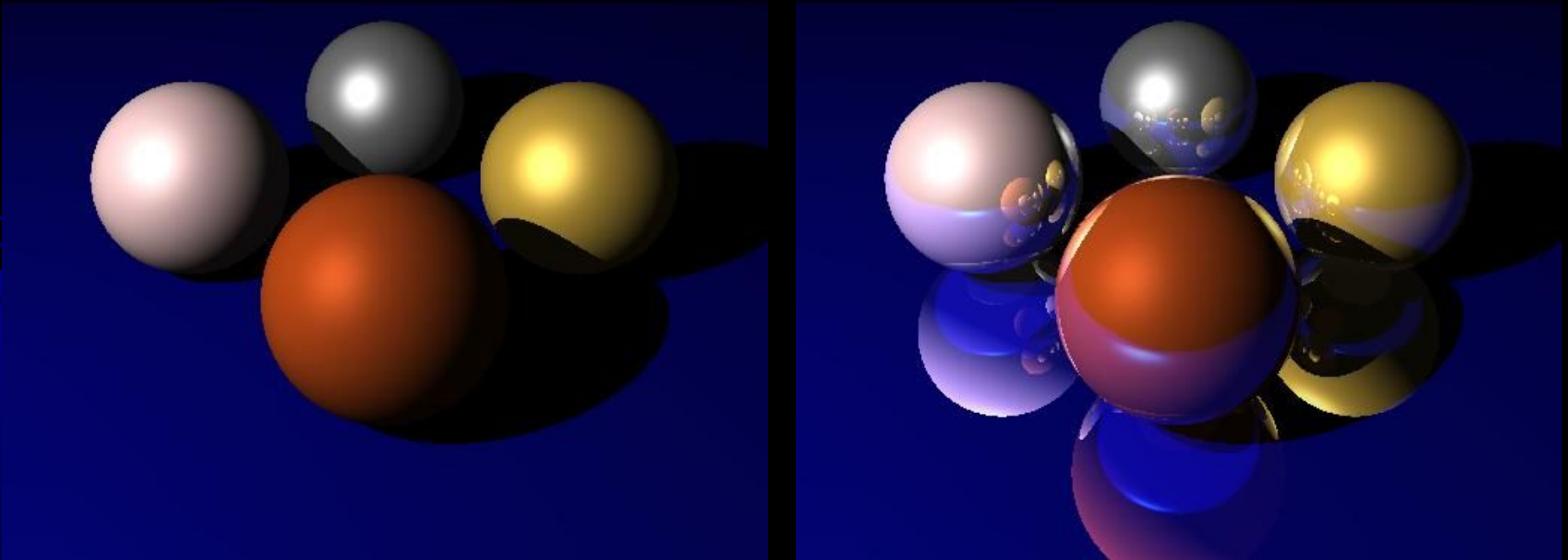


```
for each polygon
  for each scanline
    for each pixel in scanline
      update depth at pixel
      if pixel depth < buffered depth
        write to screen & depth buffers
```

# Ray Tracing

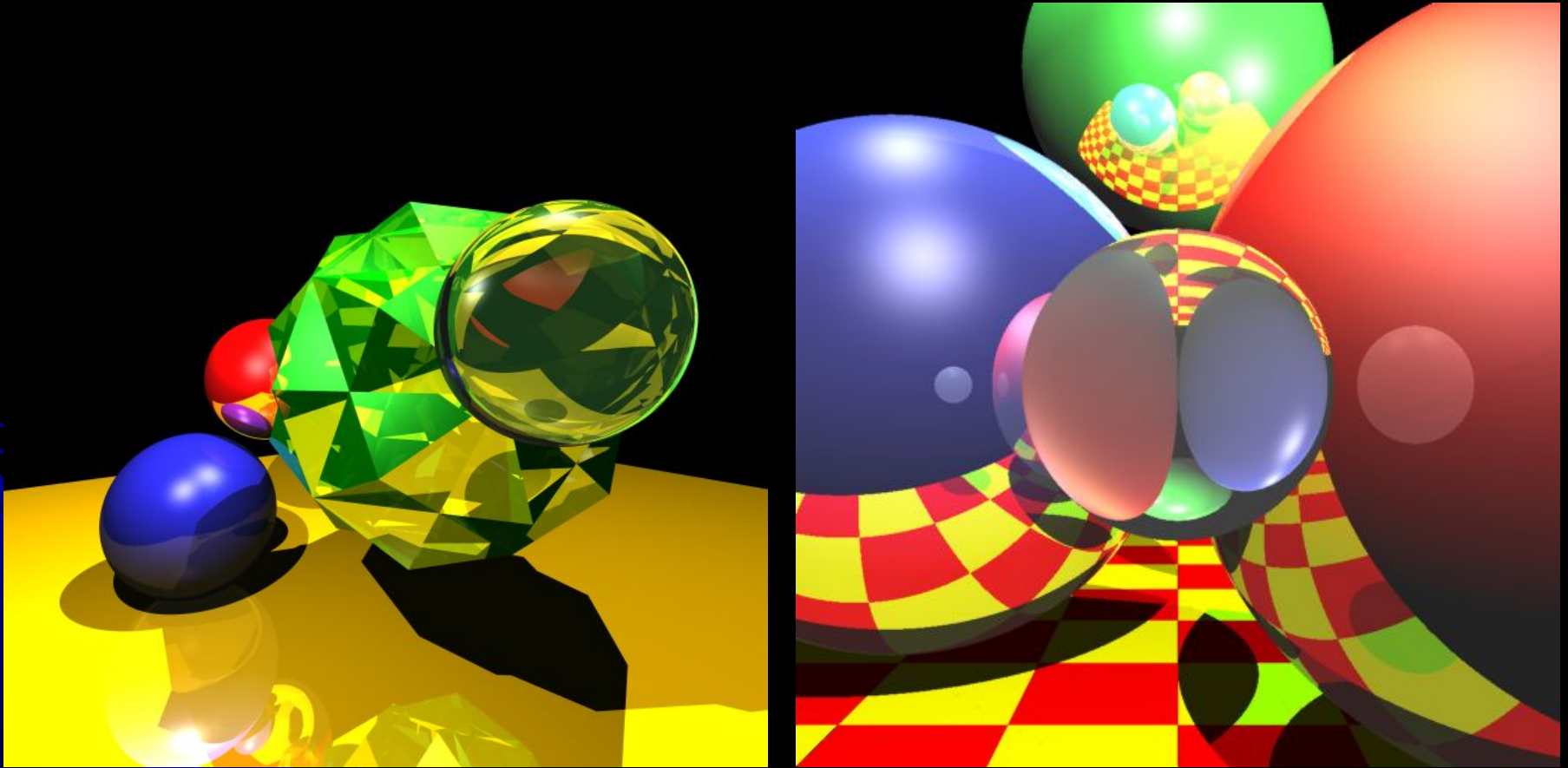
Mani Thomas  
CISC 440/640  
Computer Graphics

# Fotorealismus

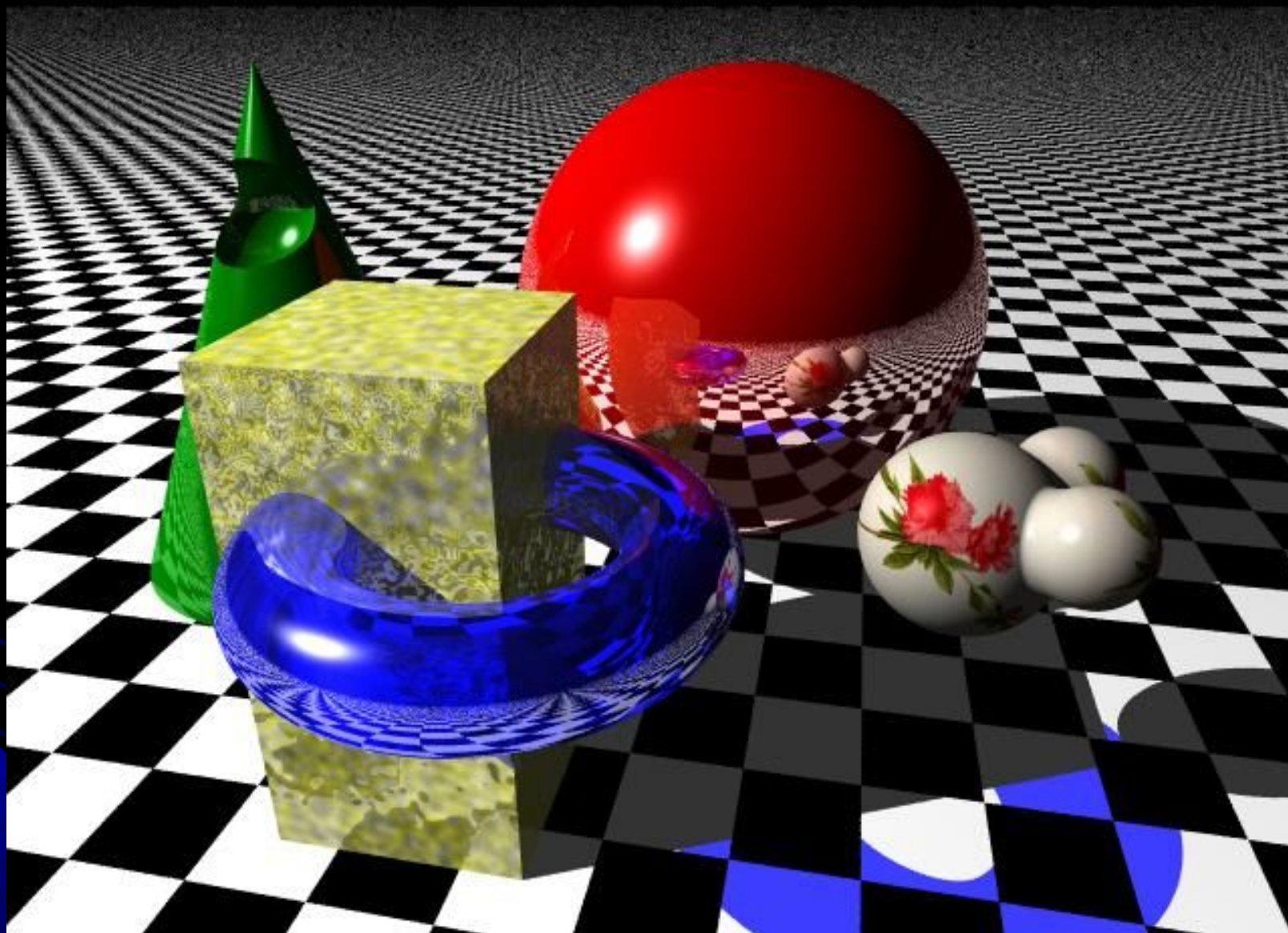


*Created by David Derman – CISC 440*

# Fotorealismus



*Created by Jan Oberlaender – CISC 640*



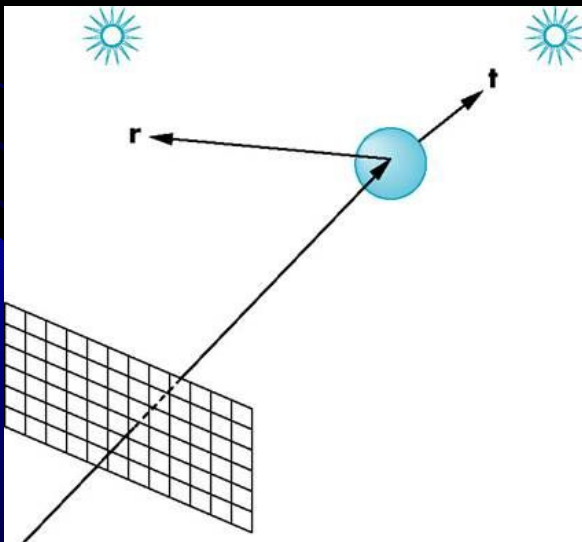
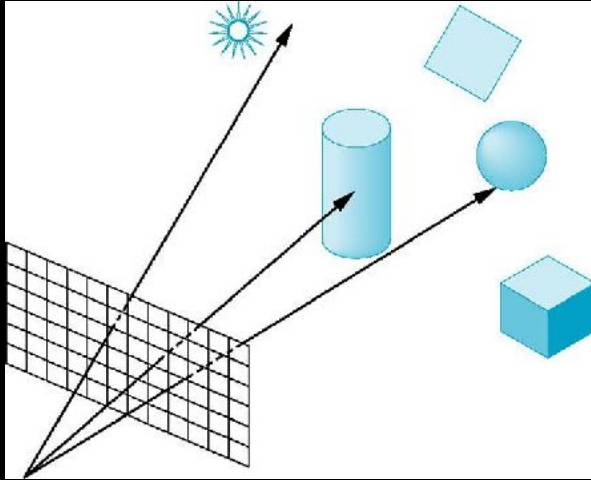
*Created by Donald Hyatt*  
*<http://www.tjhsst.edu/~dhyatt/superap/povray.html>*

# Úvod

- Co je Ray Tracing?
  - Ray Tracing je renderingová metoda založena na globálním osvětlení scény, která generuje realistické obrazy pomocí počítače.
  - V ray tracing-u, paprsek světla je sledován podél své dráhy v opačném směru.
    - Začínáme od kamery směrem ke zdroji světla a zjišťujeme stav objektů protínajících dráhu paprsku
    - Daný obrazový bod je nastaven na barvu odpovídající danému paprsku.
    - Pokud paprsek nenarazí na žádný předmět je bod nastaven na barvu pozadí.



# Ray Casting/Tracing

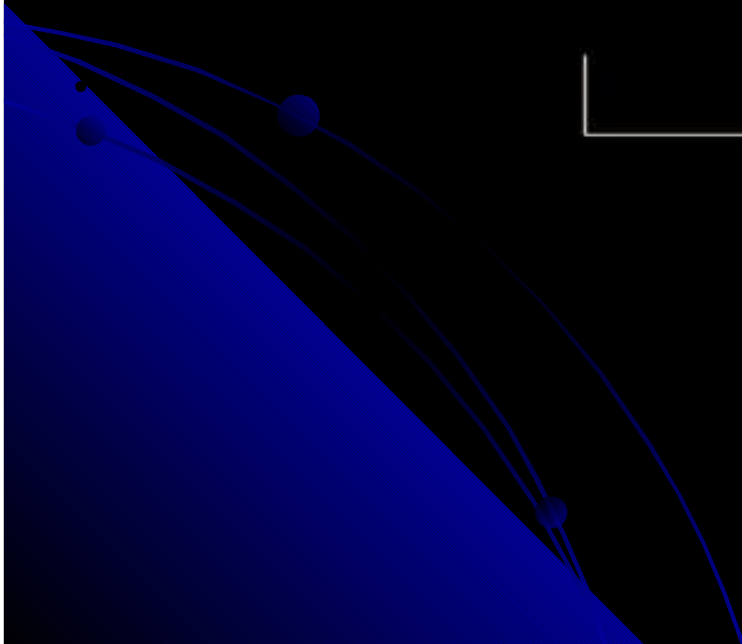
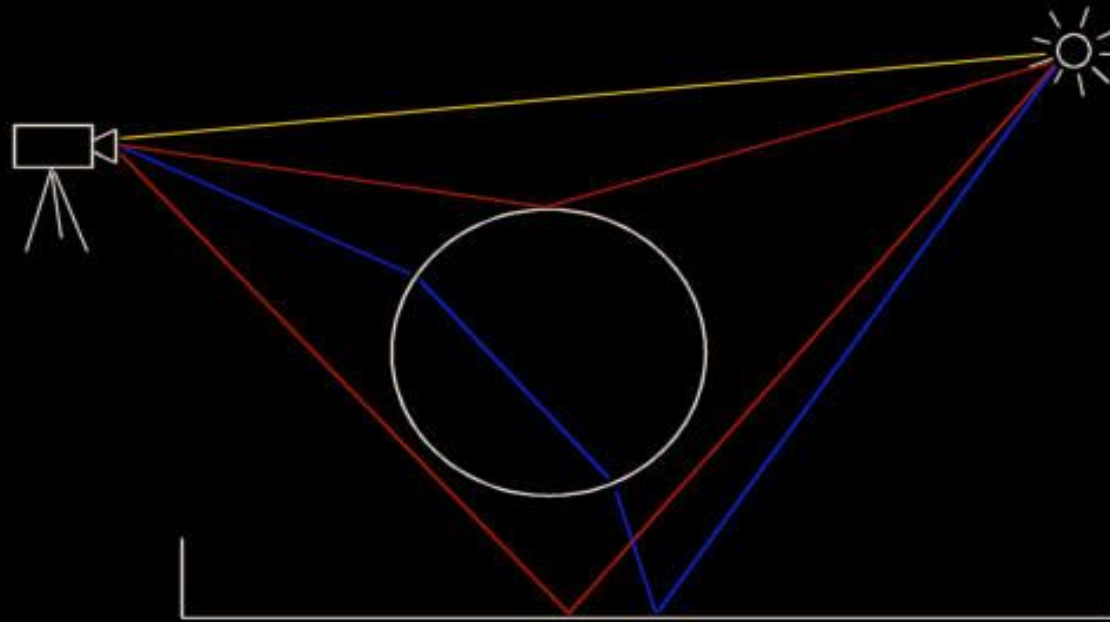


- Ray Casting
  - Paprsky se zastaví na prvním objektu
- Ray Tracing
  - Rekurze předcházejícího principu

Courtesy: Angel



# Šíření světla



# Typy paprsků

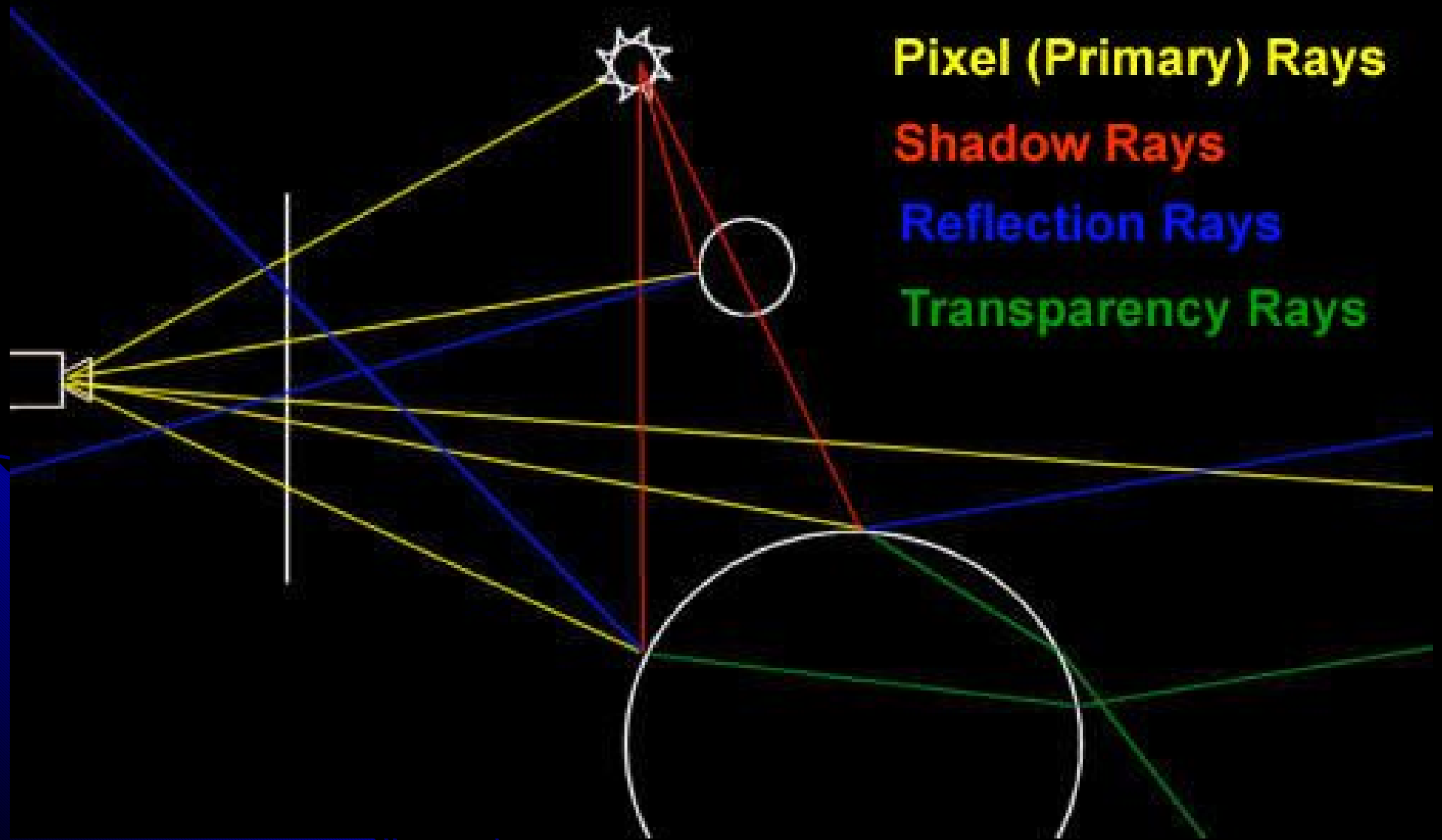


Image copyright Jacco Bikker.

# Algorithmus – Ray casting

*define the objects and light sources in the scene*

*set up the camera*

```
for(int r = 0; r < nRows; r++)
```

```
  for(int c = 0; c < nCols; c++)
```

```
  {
```

*1. Build the rc-th ray*

*2. Find all intersections of the rc-th ray with objects in the scene*

*3. Identify the intersection that lies closest to, and in front of, the eye*

*4. Compute the "hit point" where the ray hits this object, and the normal vector at that point*

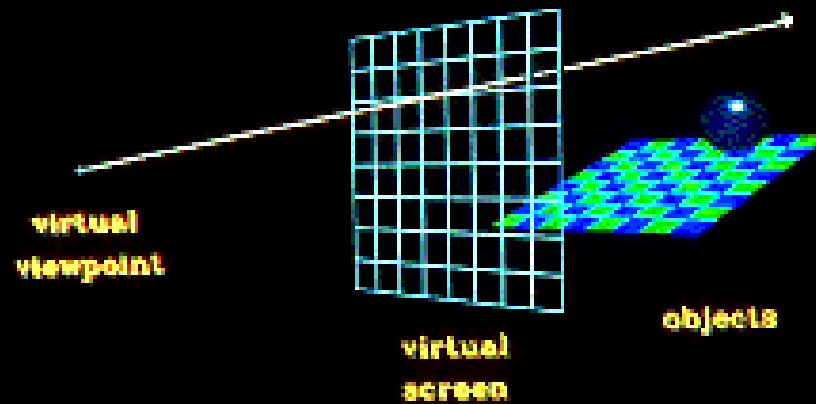
*5. Find the color of the light returning to the eye along the ray from the point of intersection*

*6. Place the color in the rc-th pixel.*

```
  }
```

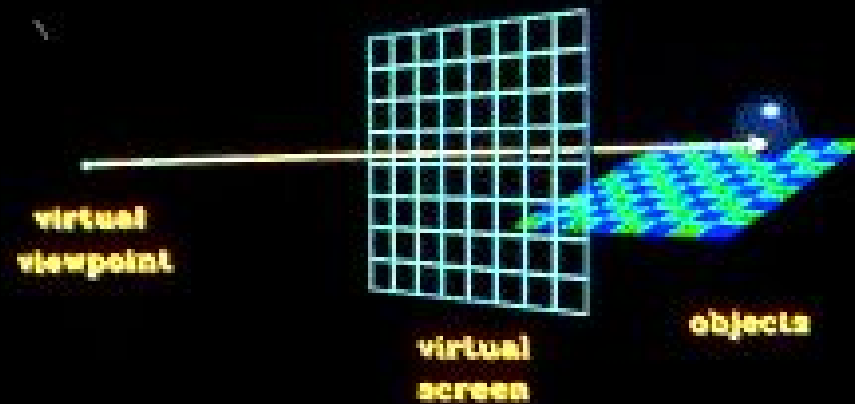
Courtesy F.S. Hill, "Computer Graphics using OpenGL"

# Ray Tracing



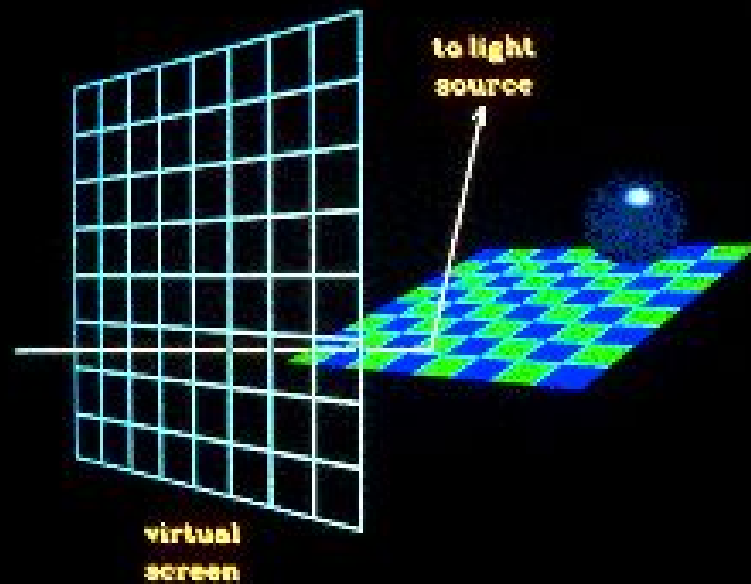
*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

# Ray Tracing



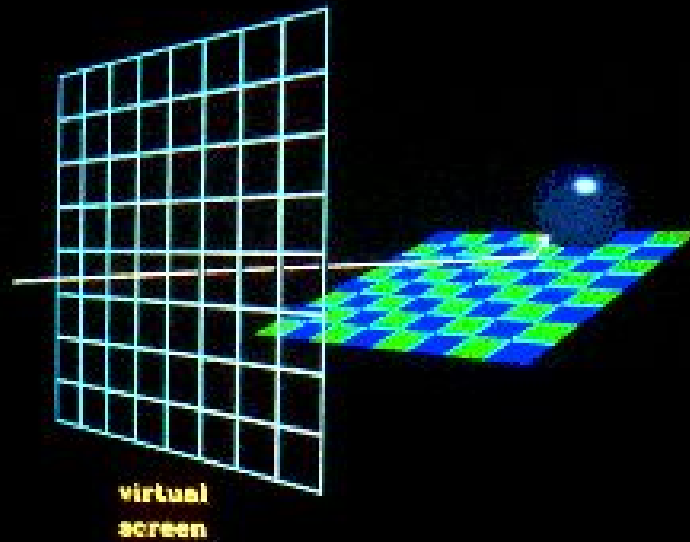
*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

# Ray Tracing



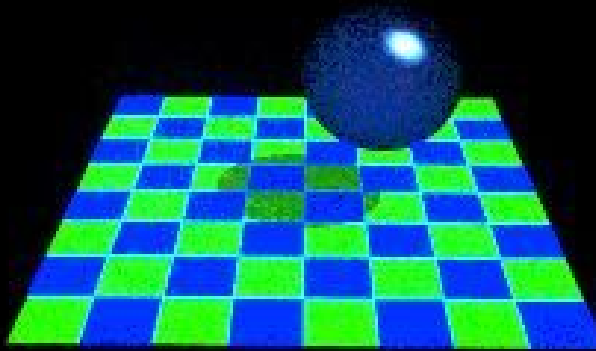
*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

# Ray Tracing



*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

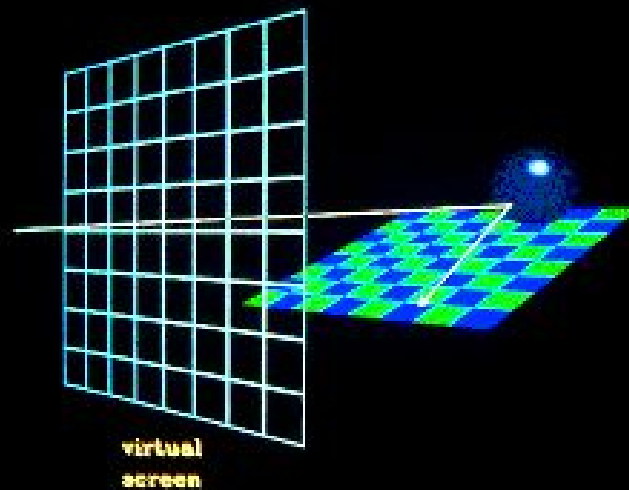
# Ray Tracing



*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

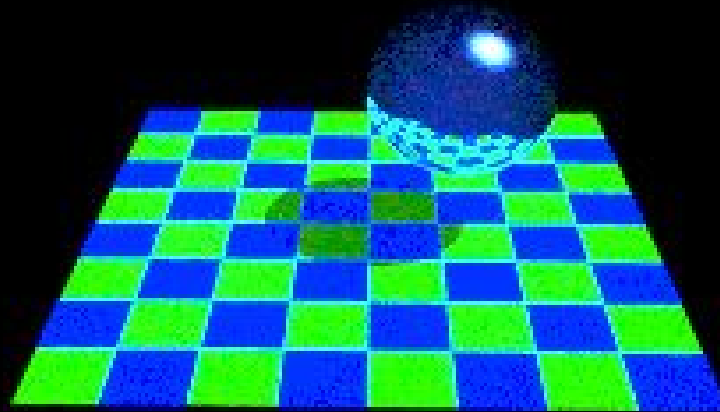


# Ray Tracing



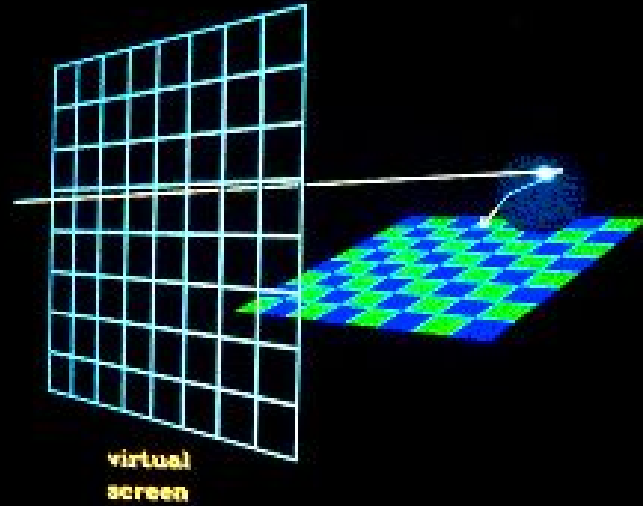
*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

# Ray Tracing



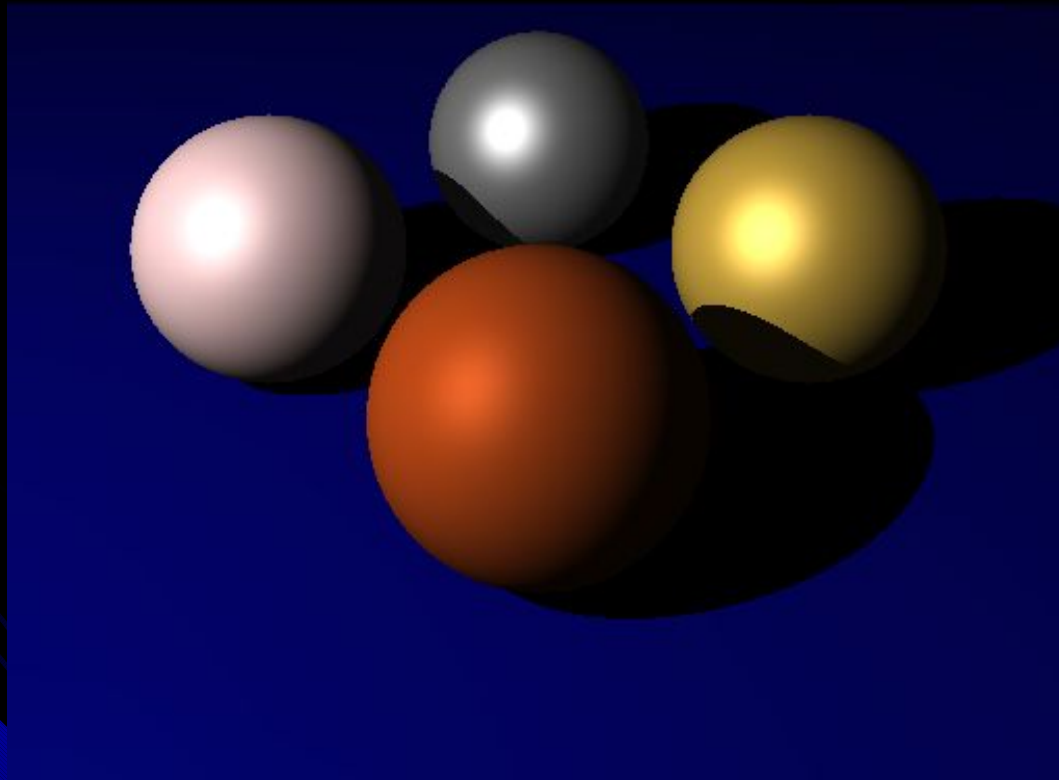
*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

# Ray Tracing

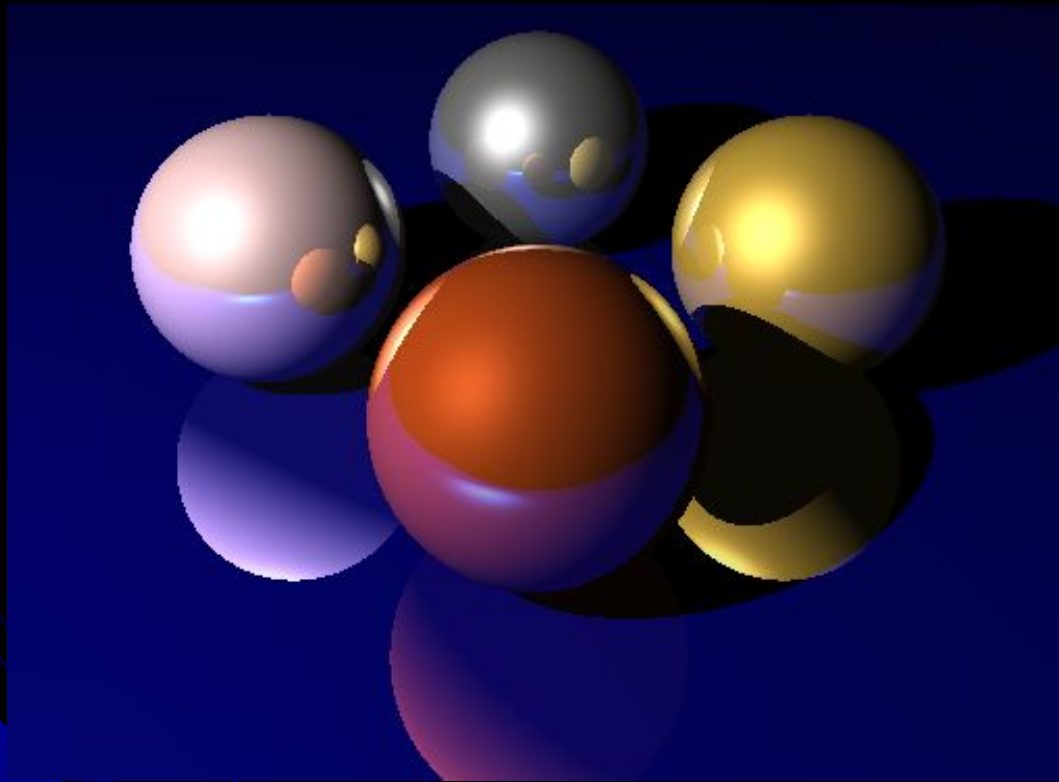


*created by Michael Sweeny, et al for ACM SIGGRAPH Education slide set 1991*

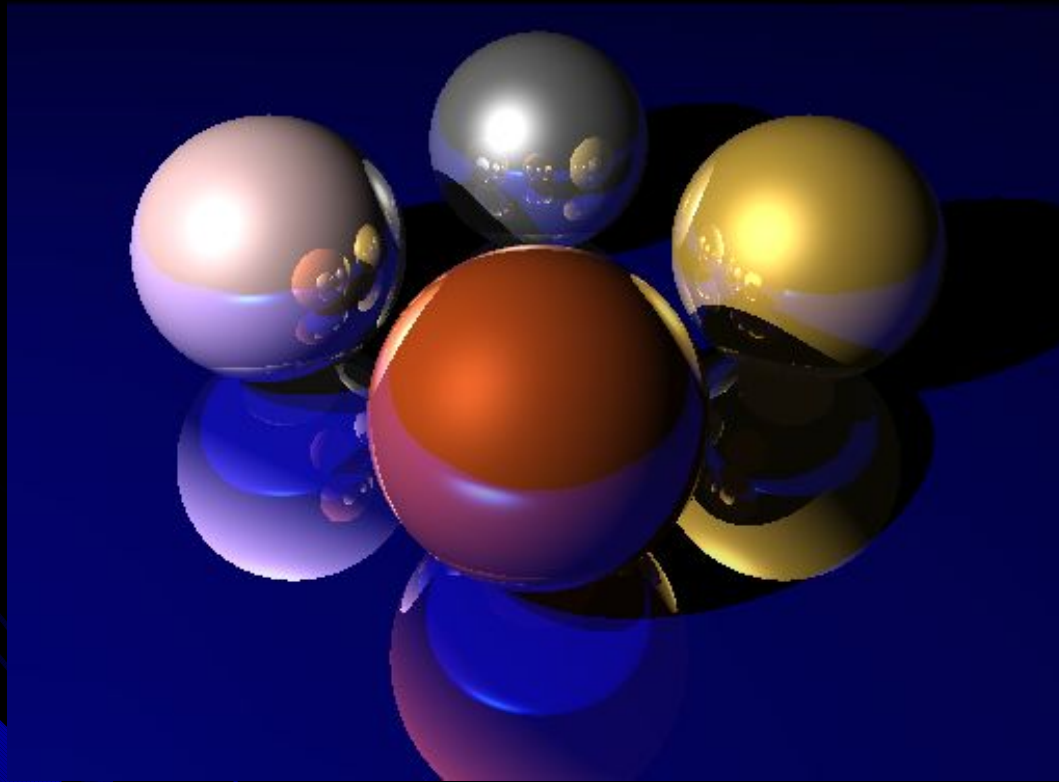
# Odraz



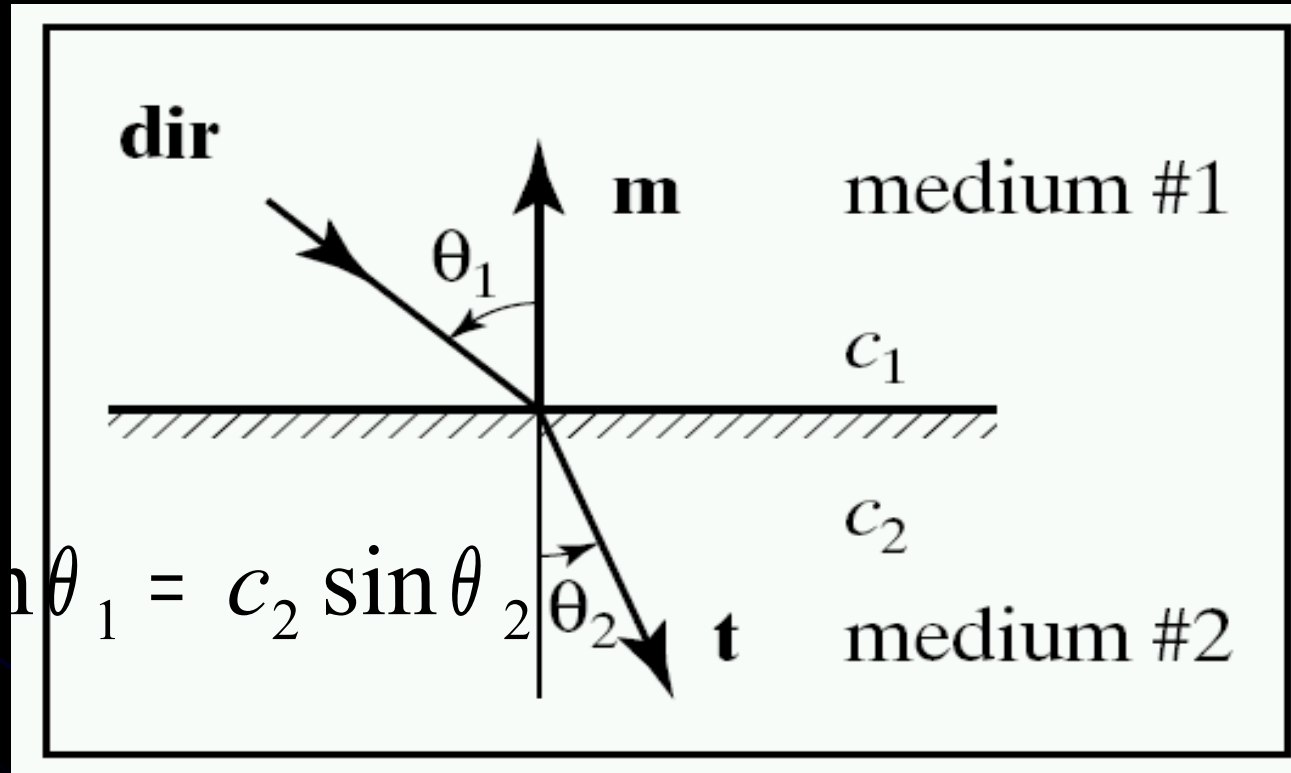
# Odraz



# Odraz



# Lom světla



Courtesy F.S. Hill, "Computer Graphics using OpenGL"

# Jiné efekty

## Hloubka ostrosti

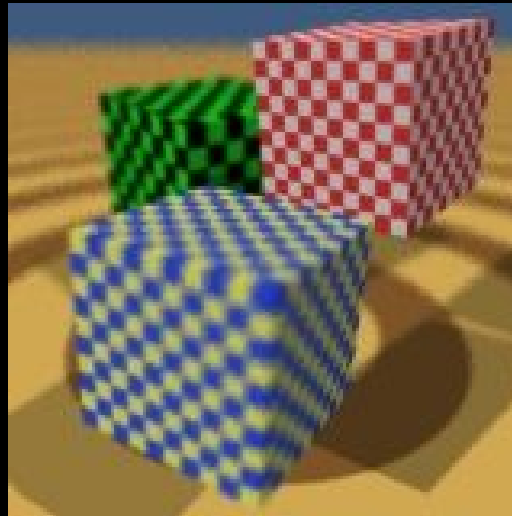


Image copyright  
Josef Pelikan  
<http://cgg.ms.mff.cuni.cz/gallery/>



# Jiné efekty

## Rozmazání pohybem

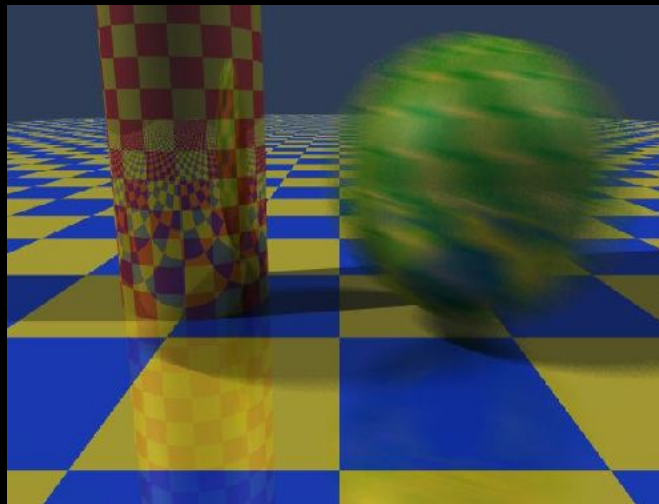
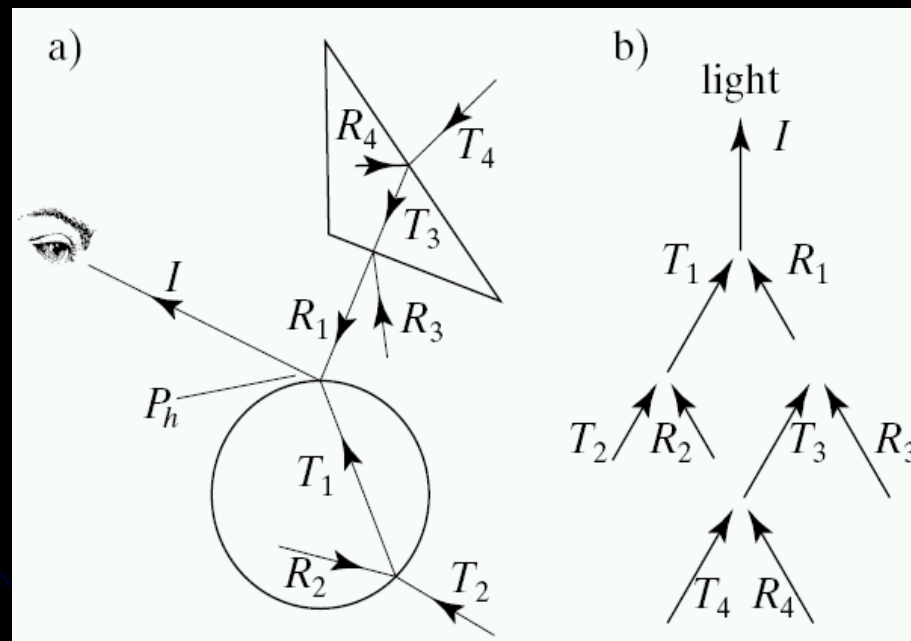


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<http://cgg.ms.mff.cuni.cz/gallery/>

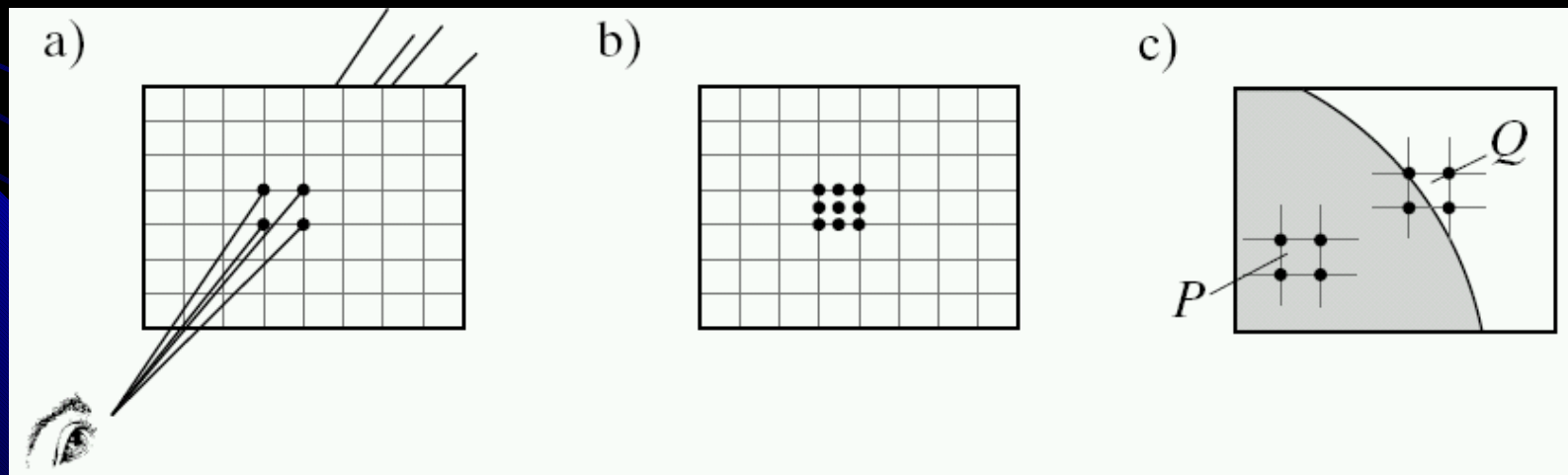
# Strom světla

- Informace o paprsku sčítají

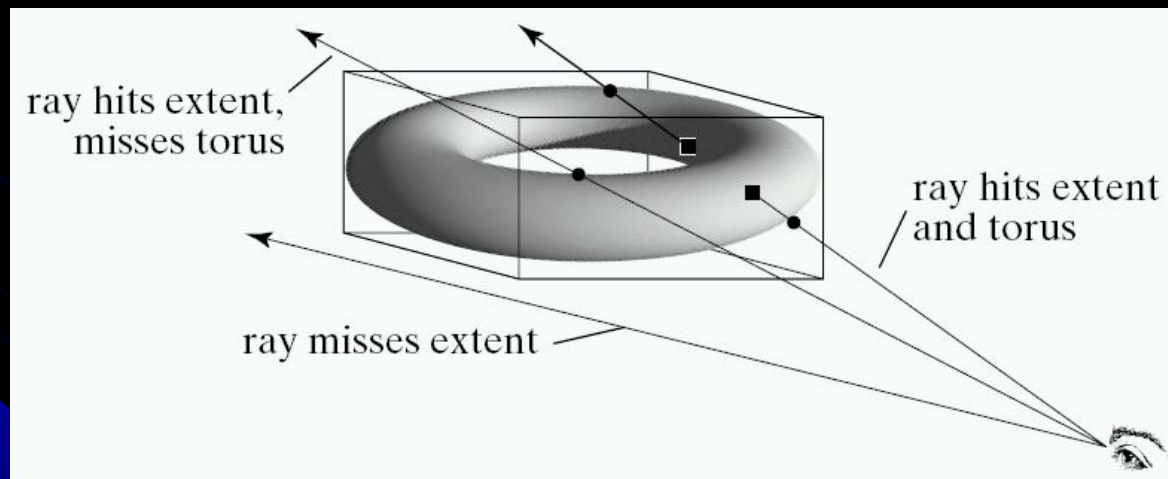


# Super-sampling

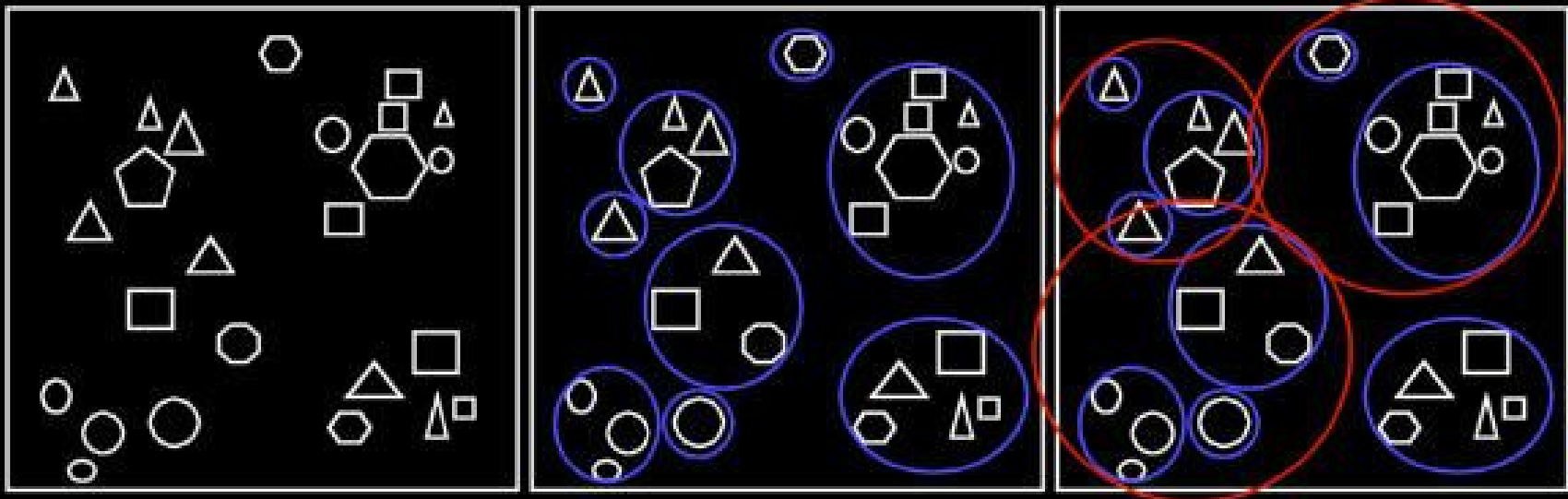
- Vyhlazení hran



# Obal



# Obal skupiny objektů





# Nerovnoměrné rozdělení na podprostory

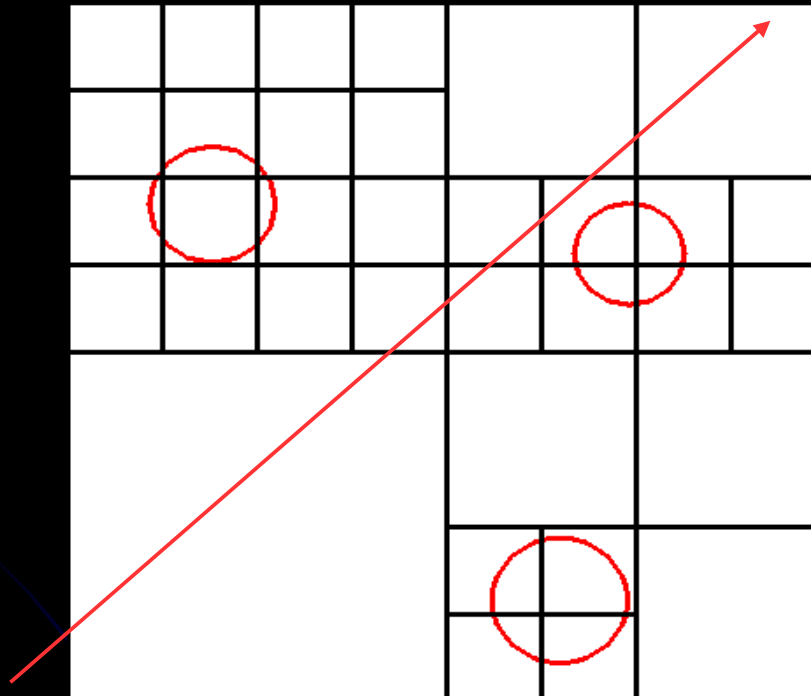


Image copyright  
Worcester Polytechnic Institute

# References

- Textbooks
  - F. S. Hill, “Computer Graphics Using OpenGL”
- Commonly used ray tracing program (completely free and available for most platforms)
  - <http://www.povray.org/>
- Interesting Links
  - Interactive Ray Tracer – Alyosha Efros
    - <http://www.cs.berkeley.edu/~efros/java/tracer/tracer.html>
- Ray Tracing explained
  - <http://www.geocities.com/jamisbuck/raytracing.html>
  - <http://www.siggraph.org/education/materials/HyperGra>



# Structure Visualization Tools

Written by James Coleman

Presented by Xiang Zhou



# Structure Visualization

- One of the primary activities in proteomics R&D is determining and Visualizing the 3D structure of proteins in order to find where drugs might modulate their activity.
- Other activities include identifying all of the proteins produced by a given cell or tissue and determining how these proteins interact.

# Some Common Tools

- 100's of visualization tools have been developed in bioinformatics.
- Many are specific to hardware such as microarray devices.
- Shareware utilities for PC's
  - PDB Viewer, WebMol, RasMol, Protein Explorer, Cn3D
  - VMD, MolMol, MidasPlus, Pymol, Chime, Chimera

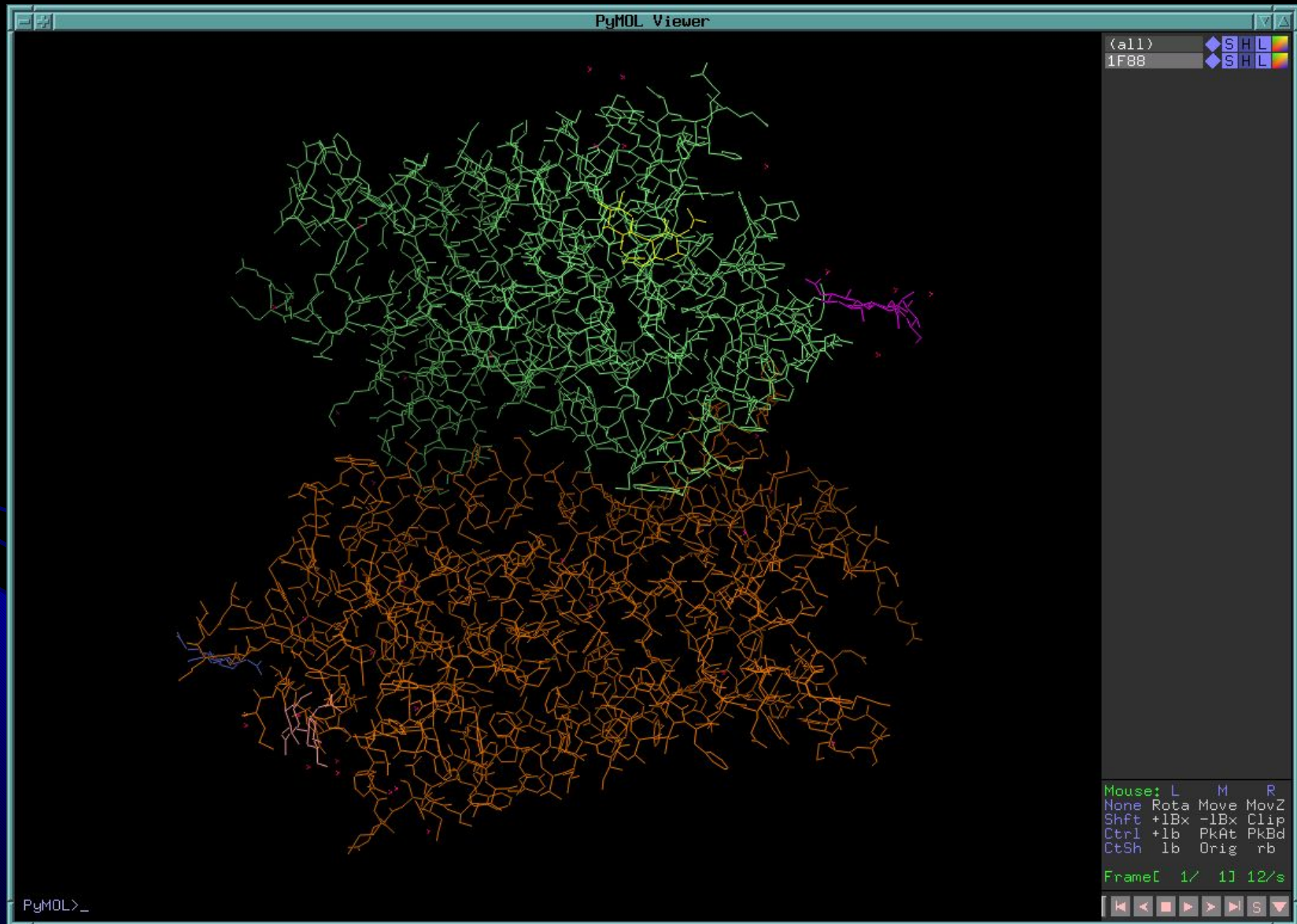
# Application Feature Summary

Feature	RasMol	Cn3D	PyMol	SWISS-PDBViewer	Chimera
<b>Architecture</b>	Stand-Alone	Plug-in	Web-Enabled	Web-enabled	Web-enabled
<b>Manipulation Power</b>	Low	High	High	High	High
<b>Hardware Requirements</b>	Low/Moderate	High	High	Moderate	High
<b>Ease of Use</b>	High; command line	Moderate	Moderate	High	Moderate;GUI +command line
<b>Special Features</b>	Small Size; easy install	Powerful GUI	GUI; ray tracing	Powerful GUI	GUI; collaboration
<b>Output Quality</b>	Moderate	Very high	High	High	Very high
<b>Documentation</b>	Good	Good	Limited	Good	Very good
<b>Support</b>	Online; Users groups	Online; Users groups	Online; Users groups	Online; Users groups	Online; Users groups
<b>Speed</b>	High	Moderate	Moderate	Moderate	Moderate/Slow
<b>OpenGL Support</b>	Yes	Yes	Yes	Yes	Yes

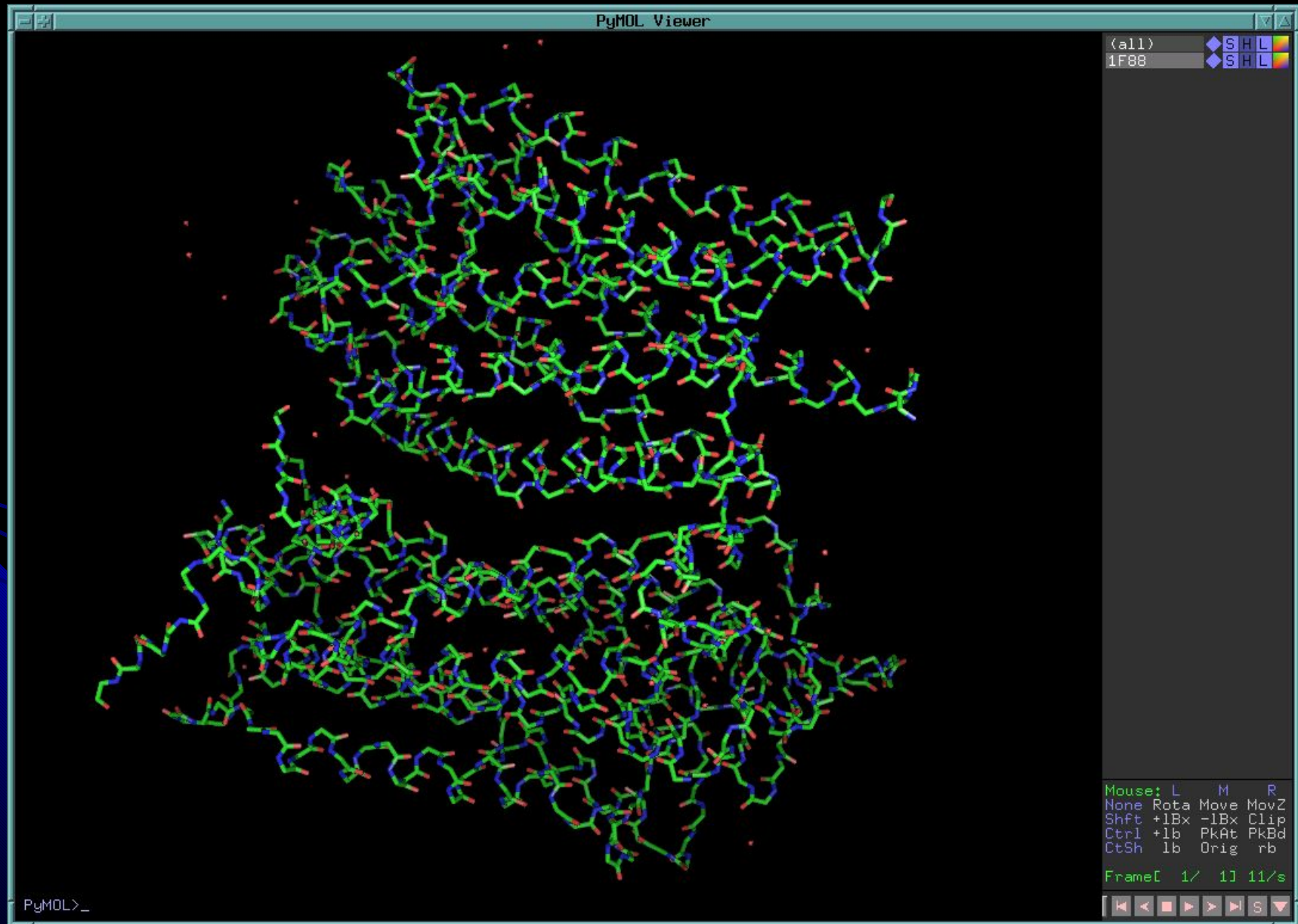
# Molecule Representations

<b>Wireframe</b>	Bonds and Bond Angles	
<b>Ball and Stick</b>	Shows Atoms, Bonds and Bonds Angles	
<b>Ribbon diagrams</b>	Shows Secondary Structure	
<b>Van der Waals surface Diagram</b>	Shows Atomic Volumes	
<b>Backbone</b>	Shows Overall Molecular Structure	

Wireframe used to show individual chains:

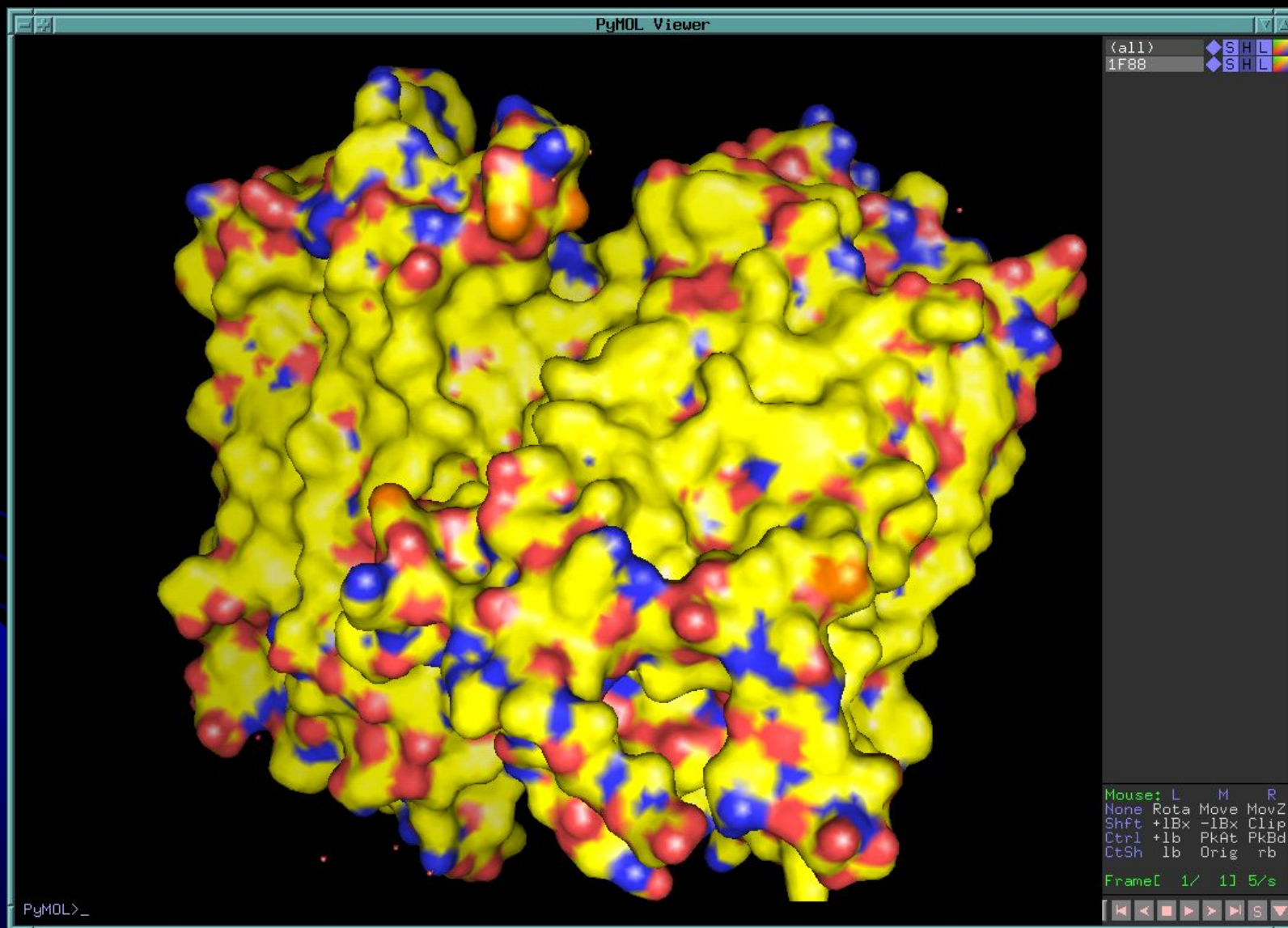


# Stick view showing atoms and bonds:



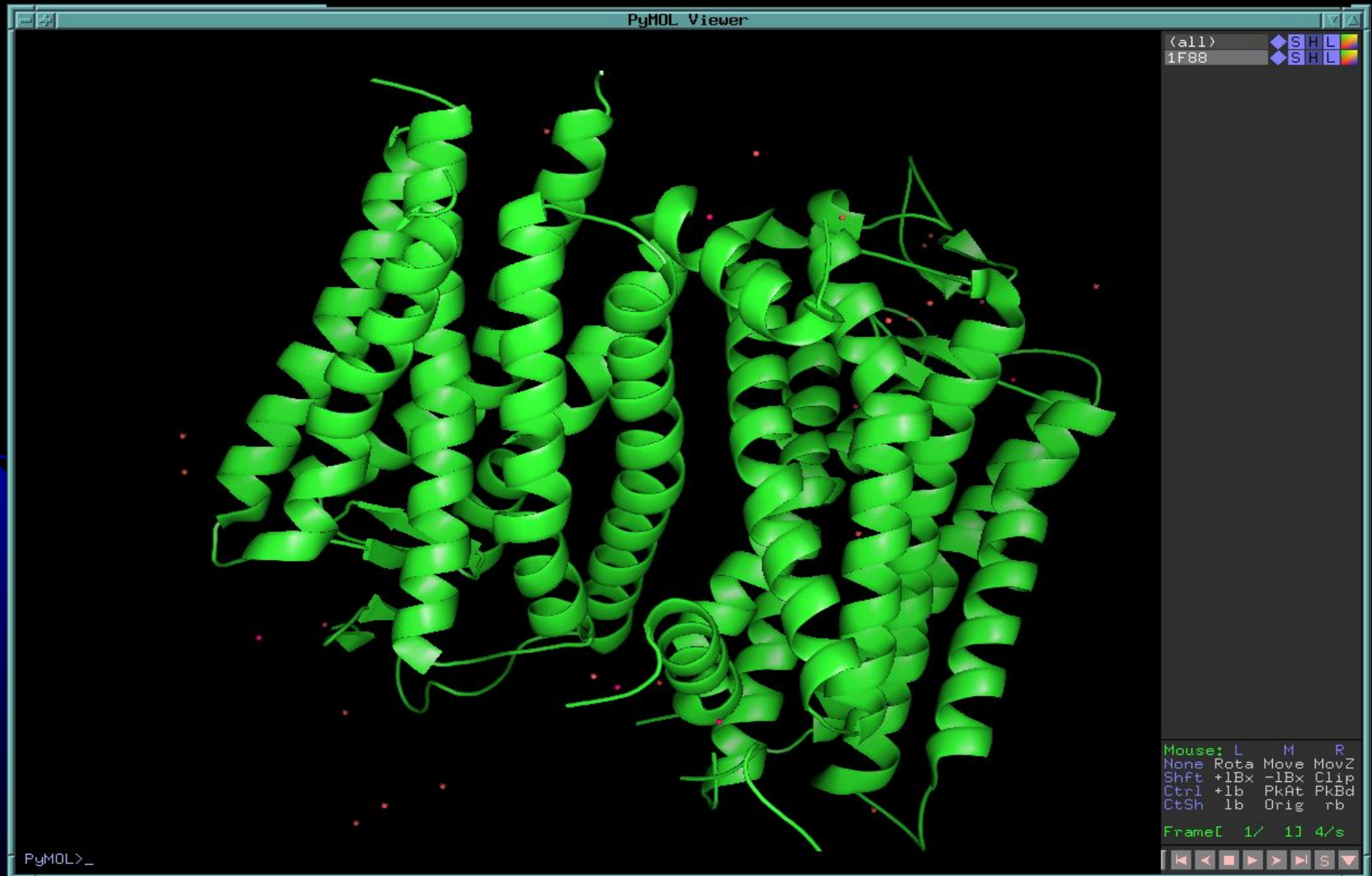


# Surface View showing surface fields:

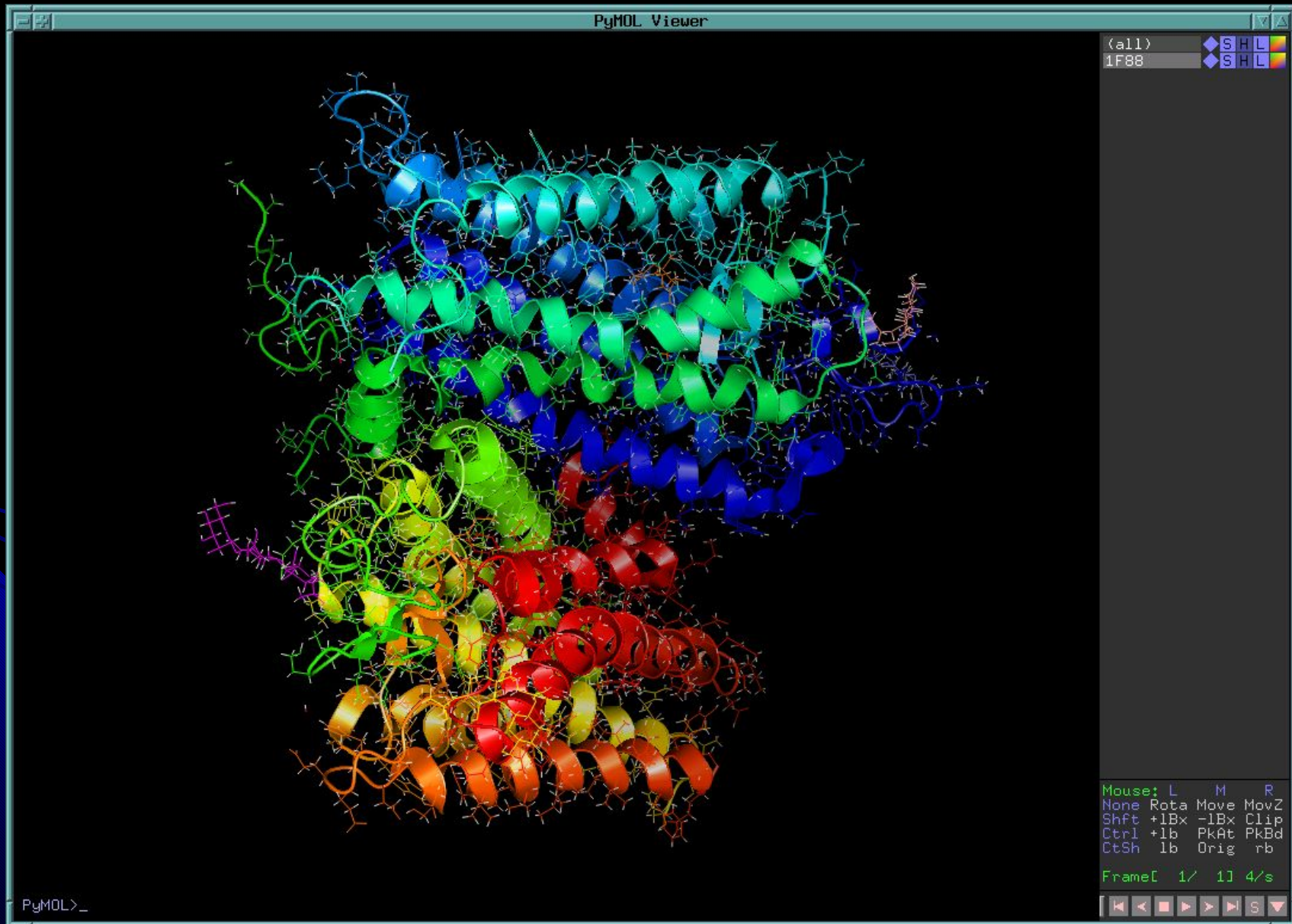




# Ribbon view of secondary structure:

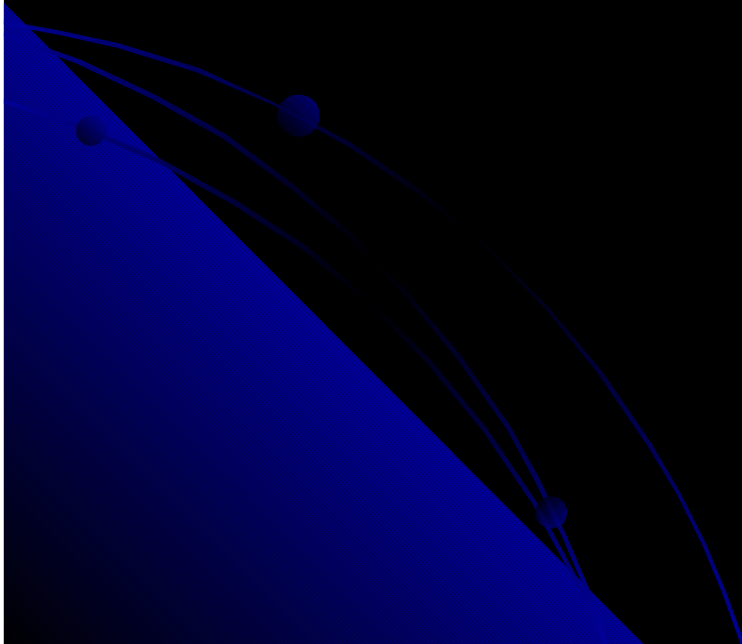


# Distinct geometrical features by color:



# Other properties that can be Visualized

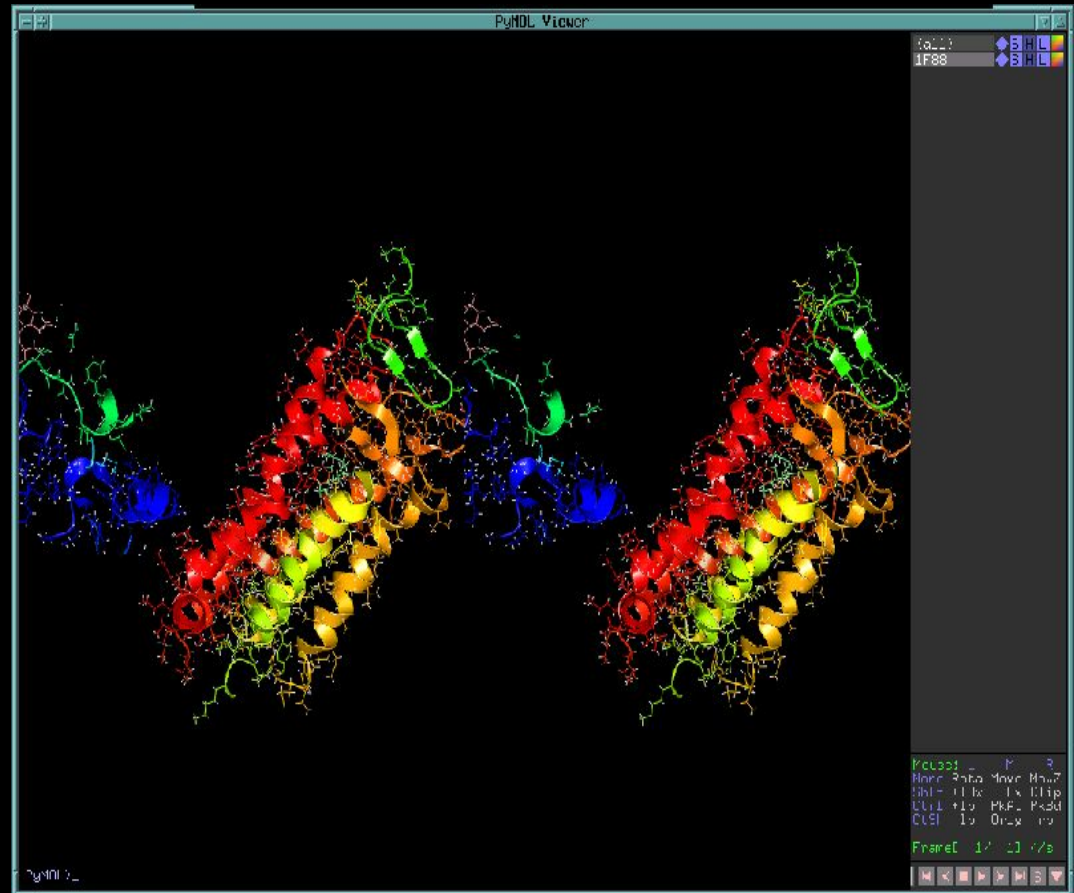
- MolMol supports the display of electrostatic potentials across a protein molecule.
- MidasPlus (a predecessor of Chimera) allows for the editing of sequences visually to see the effects of point mutations.



# For Protein interactions, we need a metaphor that reveals dynamics

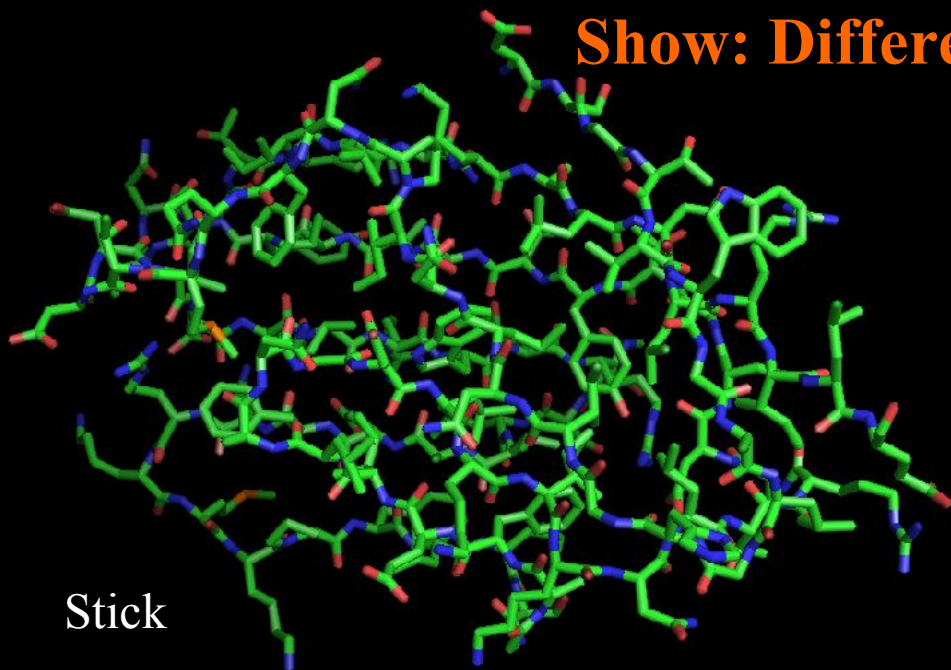
- **Haptic Joystick:** Provides force feedback when user manipulates a molecule near another one.
- **3D Goggles combined with haptic gloves** to feel electrostatic potentials and see tertiary structure dynamics.
- **PyMol provides scripting** that can produce a movie in 3D of the geometrical relationship between multiple proteins.

Stereo view of interaction of two proteins. Scripting allows for the movement of individual molecules creating a movie.

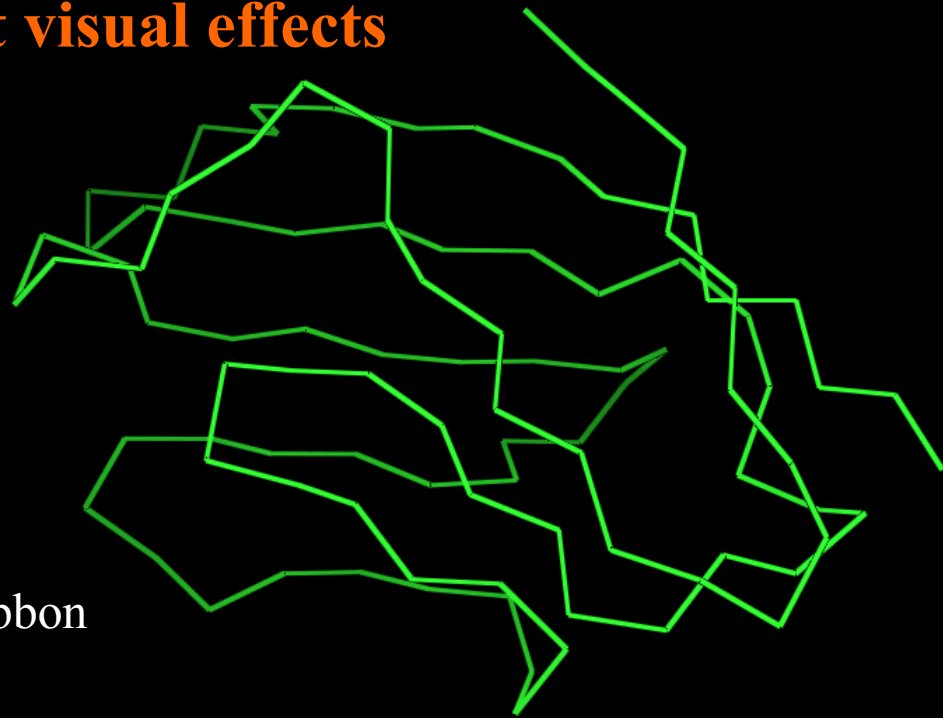




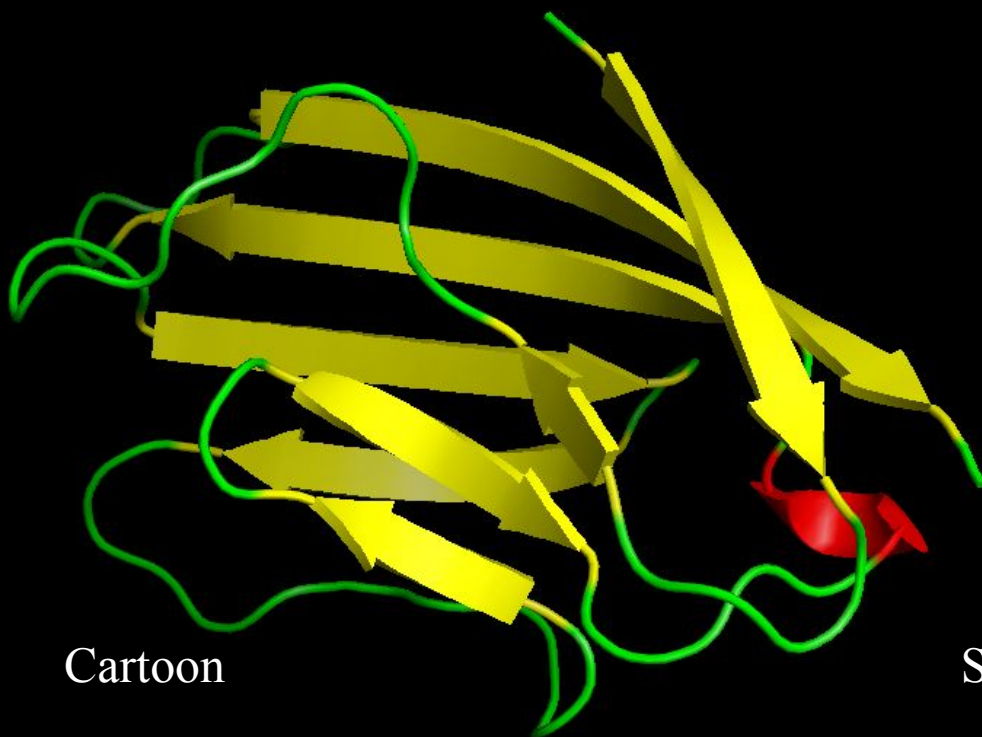
**Show: Different visual effects**



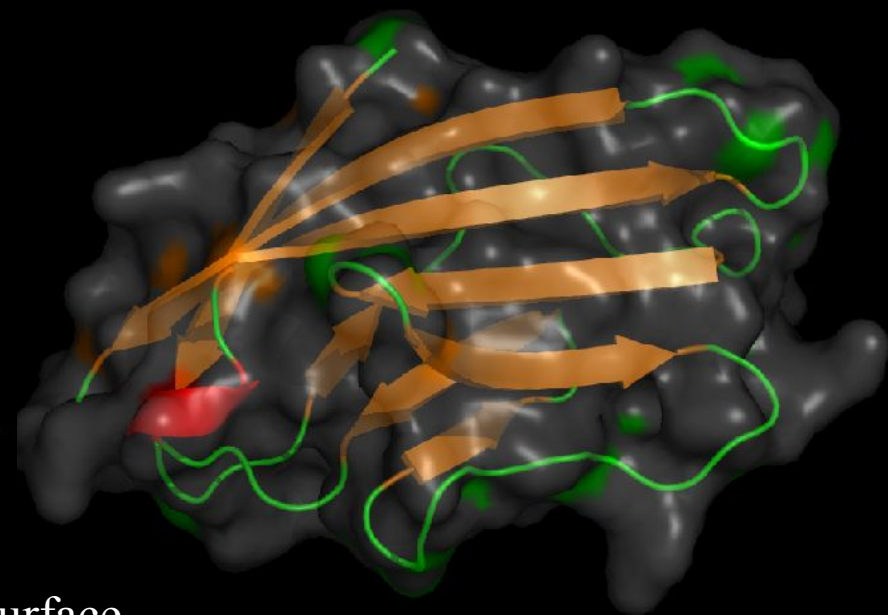
Stick



Ribbon

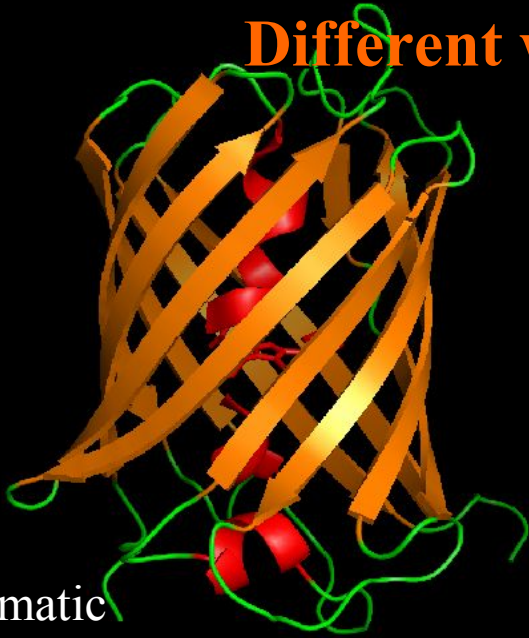


Cartoon

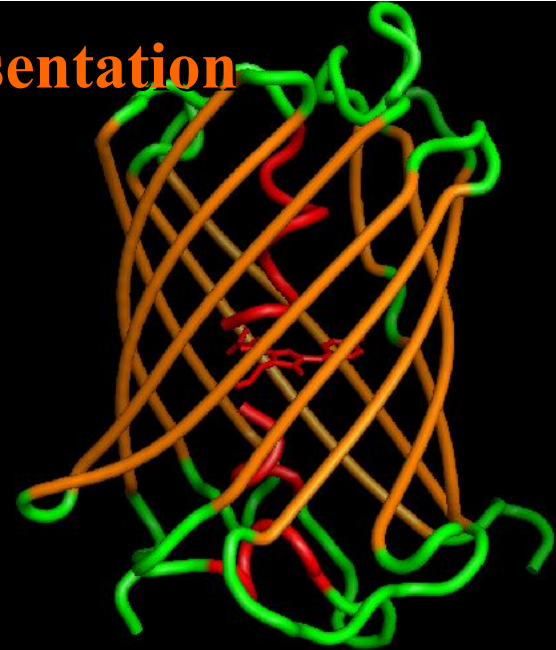


Surface

# Different way of cartoon presentation



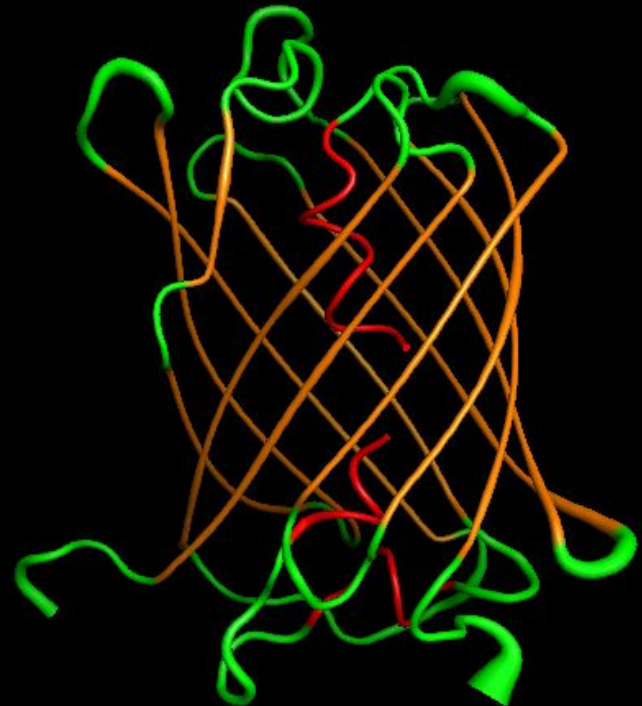
automatic



tube



loop



putty