



Convolution Primitives in HyperFun





Primitives

Algebraic primitives:

*hfSphere, hfEllipsoid, hfCylinder,
hfEllCylinder, hfEllCone, hfTorus,
hfSuperel, hfBlock*

Skeletal objects:

*hfBlobby, hfMetaball,
hfSoft*

Convolution objects:

*hfConvPoint, hfConvLine,
hfConvArc, hfConvTriangle,
hfConvCurve, hfConvMesh*

Procedural objects:

hfNoiseG

Operations

*hfScale,
hfShift,
hfRotate,
hfTwist,
hfStretch,
hfTaper
hfBlendUni,
hfBlendInt*

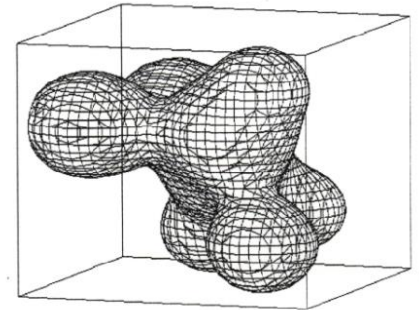
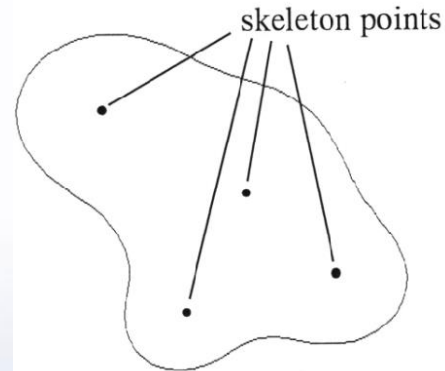


Skeletal Surface Definition

$$F(P) - T = 0$$

with

$$F(P) = \sum_{i=1}^N c_i F_i(r_i)$$



N is the number of skeletal elements,

F_i is the individual scalar field,
(*blending function*) of the i -th element,

r_i is the distance from P to the i -th element,

T is the *threshold* (or *level value*).



Convolution Integral

Defining function of a convolution primitive:

$$f(X) = \int_{R^3} s(P)h(X - P)dP$$

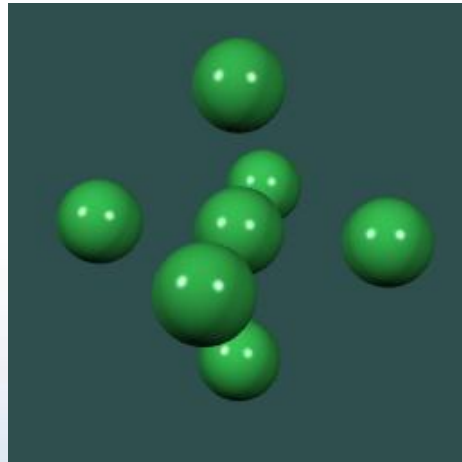
- $s(X)$ is a predicate function defining geometry of the skeletal element
- $h(X)$ is a convolution kernel

Usually integration requires heavy numerical calculations, but we use analytical solutions for integrals over several skeletal elements.

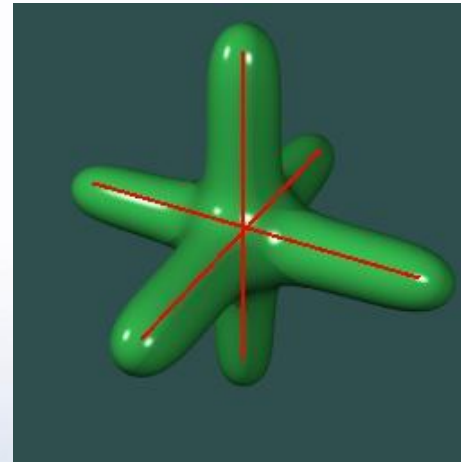


Skeletal elements

Point

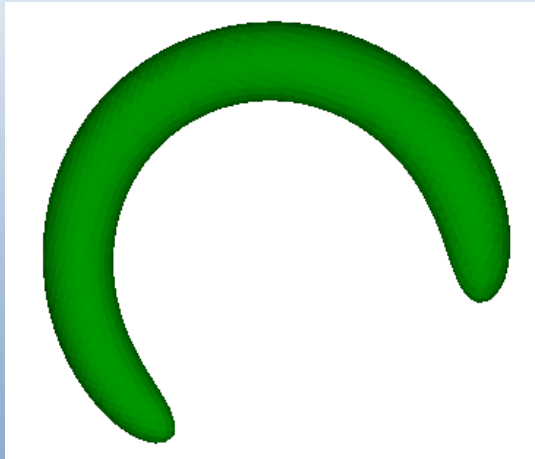


Line

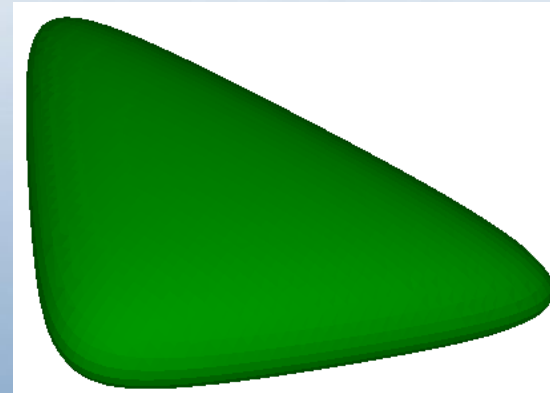


Images by Yuichiro Goto

Arc



Triangle





Convolution primitive: Skeletal Points

hfConvPoint(x,vect,S,T)

- **x** – given point coordinates for the function evaluation;
- **vect** – linear array of skeleton points' coordinates organized as $(x_1, y_1, z_1, x_2, y_2, z_2, \dots)$;
- **S** - array of inverse kernel width parameters for each skeletal point; smaller S_i means bigger i-th component;
- **T** - threshold value for the entire model; smaller **T** means entire expanded surface; bigger **T** means entire contracted surface.

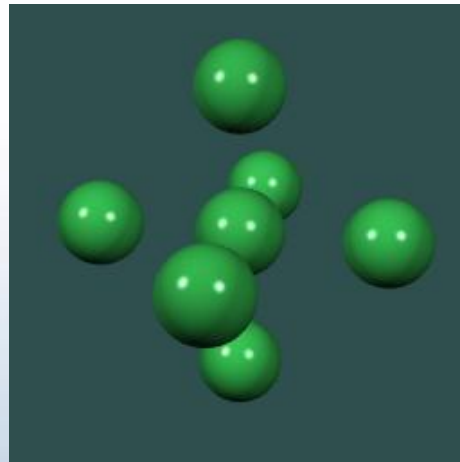


Convolution primitive: Skeletal Points

Smaller S_i means bigger i -th component

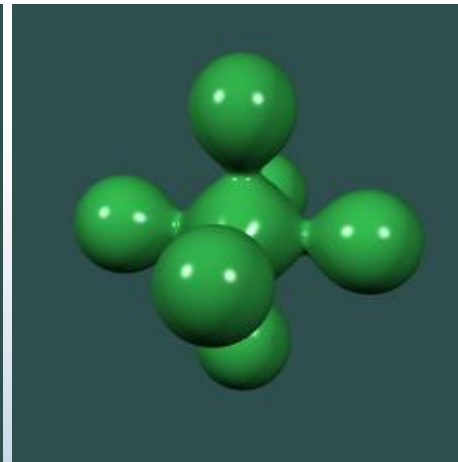
$S = 1.0$

$T = 0.1$



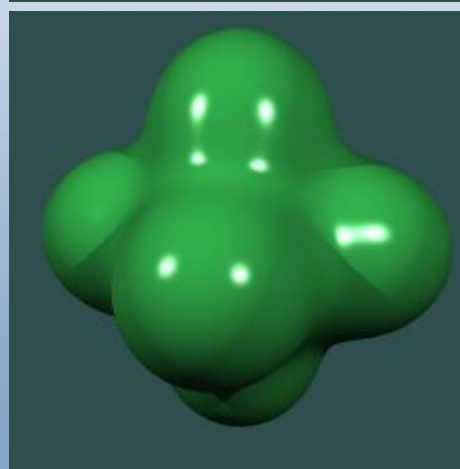
$S = 0.75$

$T = 0.1$



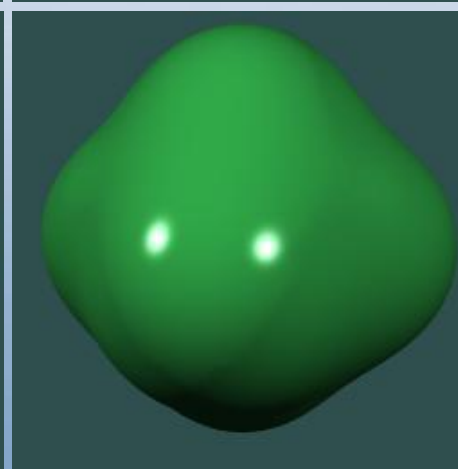
$S = 0.5$

$T = 0.1$



$S = 0.35$

$T = 0.1$





Convolution primitive: Skeletal Lines

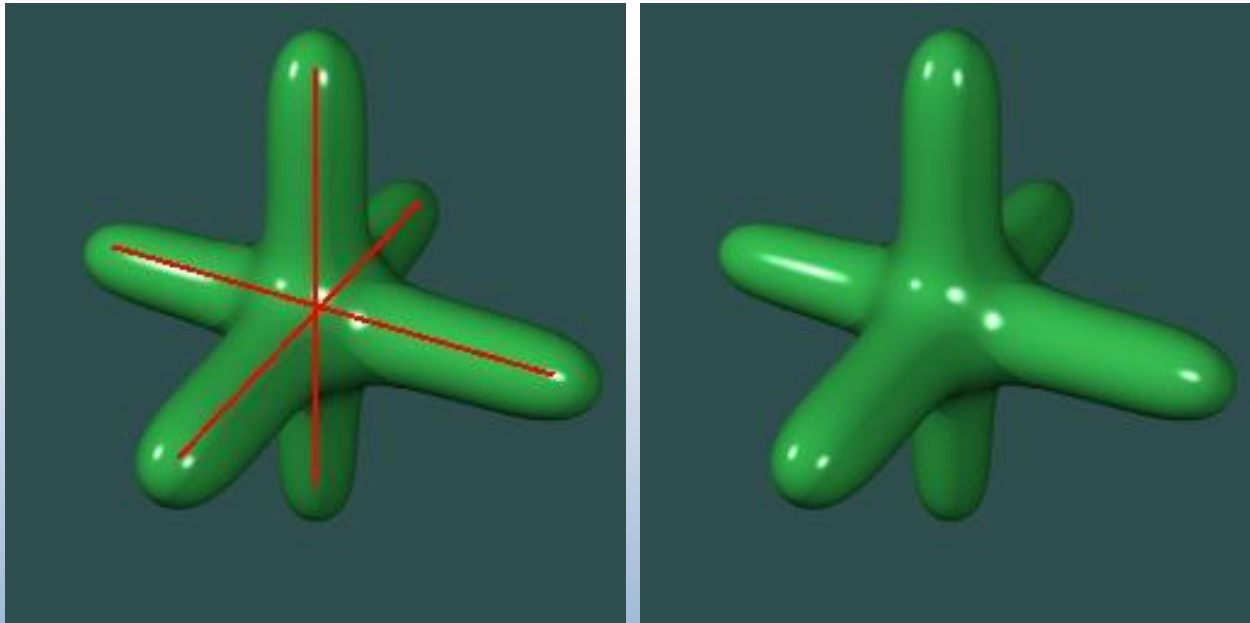
hfConvLine(x,begin,end,S,T)

- **x** – given point coordinates for the function evaluation;
- **begin** – linear array of beginning points' coordinates of line segments, organized as $(x_{b1}, y_{b1}, z_{b1}, x_{b2}, y_{b2}, z_{b2}, \dots)$;
- **end** – array of ending points' coordinates of line segments, organized as $(x_{e1}, y_{e1}, z_{e1}, x_{e2}, y_{e2}, z_{e2}, \dots)$;
- **S** - array of inverse kernel width parameters for each skeletal line segment; smaller S_i means bigger i-th component;
- **T** - threshold value for the entire model; smaller **T** means entire expanded surface; bigger **T** means entire contracted surface.



Convolution primitive: Skeletal Lines

Convolution primitive defined by three line segments.



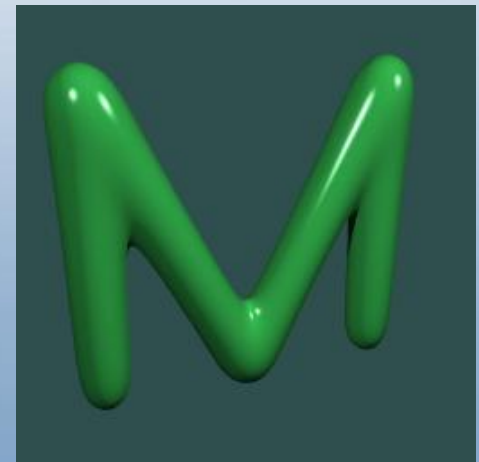
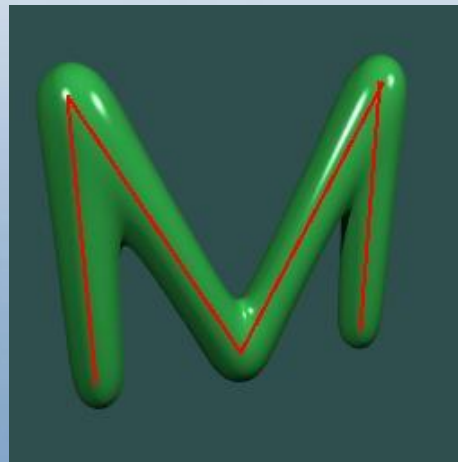


Convolution primitive: Skeletal Curve

hfConvCurve(x,vect,S,T)

- **x** – given point coordinates;
- **vect** – linear array of skeleton curve points' coordinates organized as $(x_1, y_1, z_1, x_2, y_2, z_2, \dots)$;
- **S** - array of inverse kernel width parameters;
- **T** - threshold.

Convolution surface
with a skeleton curve
defined by five points.



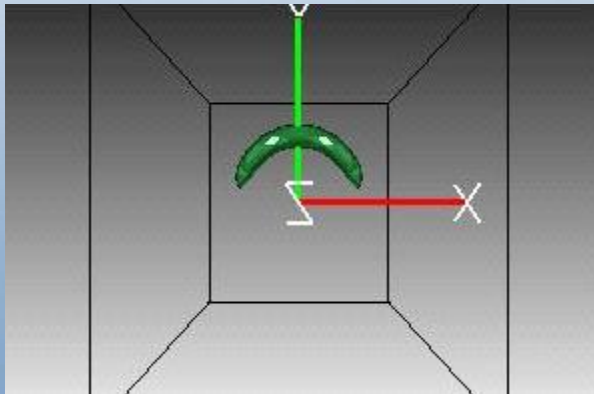


Convolution primitive: Skeletal Arcs

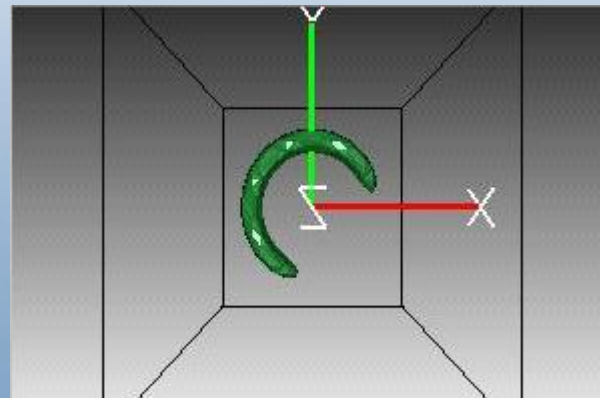
hfConvArc(x,center,radius,theta,axis,angle,S,T)

- **x** – given point coordinates
- **center** – coordinate array for centers of arcs
- **radius** – array of arcs' radii
- **theta** – array of arcs' angles measured from positive x-axis counter-clockwise, 360 degrees are used for the full circle)
- **axis** – array of vectors defining axis of rotation for each arc placed on a local plane parallel to the xy-plane
- **angle** – angles of rotation for arcs around axis of rotation
- **S** - array of inverse kernel width parameters
- **T** - threshold.

theta = 180



theta = 270





Convolution primitive: Skeletal Arcs

- Convolution primitive defined by two skeletal arcs
- two full circles with $\theta = 360$
 - one rotated about x-axis

```
arcs(x[3], a[1]) {
```

```
.....
```

```
theta = [ 360.0, 360.0];
```

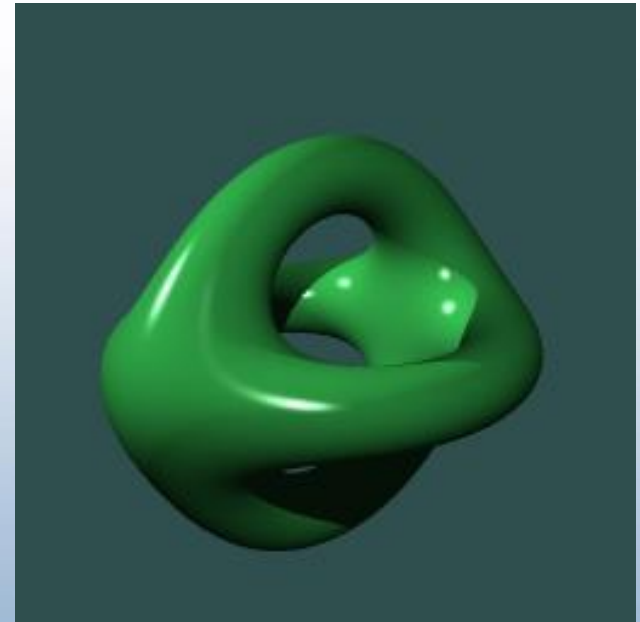
```
axis = [ 0.0, 0.0, 1.0,  
        1.0, 0.0, 0.0];
```

```
angle = [ 0.0, 90.0];
```

```
s = [ 0.5, 0.5 ];
```

```
arcs = hfConvArc(x, center, radius, theta, axis,  
                angle, s, 0.5);
```

```
}
```



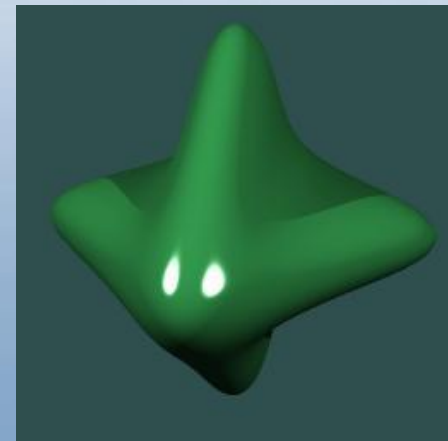


Convolution primitive: Skeletal Triangles

hfConvTriangle(x,vect,S,T)

- **x** – given point coordinates;
- **vect** – coordinate array for vertices of triangles, 9 elements for each triangle organized as $(x_1, y_1, z_1, x_2, y_2, z_2, x_3, y_3, z_3 \dots)$;
- **S** - array of inverse kernel width parameters;
- **T** - threshold.

Convolution surface with
four skeleton triangles.





Convolution primitive: Skeletal Mesh

hfConvMesh(x,vertex,index,S,T)

- **x** – given point coordinates;
- **vertex** – coordinate array for vertices of connected triangles organized as $(x_1, y_1, z_1, x_2, y_2, z_2, x_3, y_3, z_3 \dots)$;
- **index** – list of vertex indices, 3 per triangle organized as (i_1, i_2, i_3, \dots)
- **S** - array of inverse kernel width parameters;
- **T** - threshold.

```
vertex = [  
-2.5, 0.0, 0.0,  
0.0, 2.5, 0.0,  
2.5, 0.0, 0.0,  
0.0, -2.5, 0.0];  
index = [ 1, 2, 3, 1, 4, 3 ];
```

Two triangles
described in
vertex and **index**
arrays – memory
saving structure