```
at a = nt - nc, b
(AXDEPTH)
survive = SurvivalProbability( diff.
e.x + radiance.y + radiance.z) > 0)
v = true;
at brdfPdf = EvaluateDiffuse( L, N )
at3 factor = diffuse * INVPI;
at weight = Mis2( directPdf, brdfPdf );
at cosThetaOut = dot( N, L );
E * ((weight * cosThetaOut) / directPdf)
andom walk - done properly, closely followi
at3 brdf = SampleDiffuse( diffuse, N, r1, r2, &R, Apo
ırvive;
n = E * brdf * (dot( N, R ) / pdf);
```

Ray Tracing for Games

Dr. Jacco Bikker - IGAD/BUAS, Breda, February 3

Welcome!



- Combining BVHs
- Refitting
- Rigid Motion



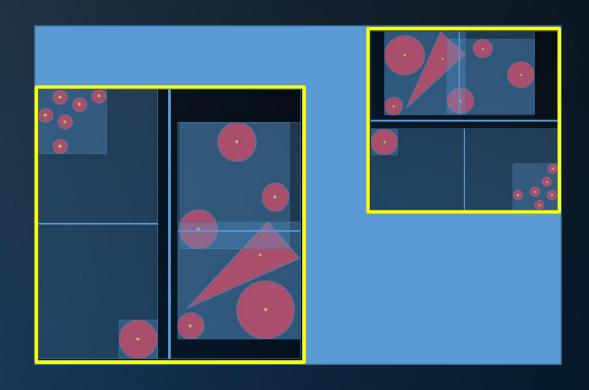






Combining BVHs

```
(AXDEPTH)
survive = SurvivalProbability( diffus
radiance = SampleLight( &rand, I, &L, &L
e.x + radiance.y + radiance.z) > 0) 88
v = true;
at brdfPdf = EvaluateDiffuse( L, N ) * Ps
at3 factor = diffuse * INVPI;
at weight = Mis2( directPdf, brdfPdf );
at cosThetaOut = dot( N, L );
E * ((weight * cosThetaOut) / directPdf) * (rad)
/ive)
at3 brdf = SampleDiffuse( diffuse, N, r1, r2, &R, &pdf
ırvive;
pdf;
n = E * brdf * (dot( N, R ) / pdf);
```





Ray Tracing for Games

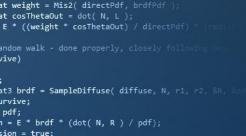
Combining BVHs

```
BVH #1: Car 1
BVH #2: Car 2
BVH #3: Track
```

Per frame:

- Rebuild BVH for each car
- Combine car 1 and car 2
- Combine track and cars
- Trace.





(AXDEPTH)

v = true;

survive = SurvivalProbability(diff

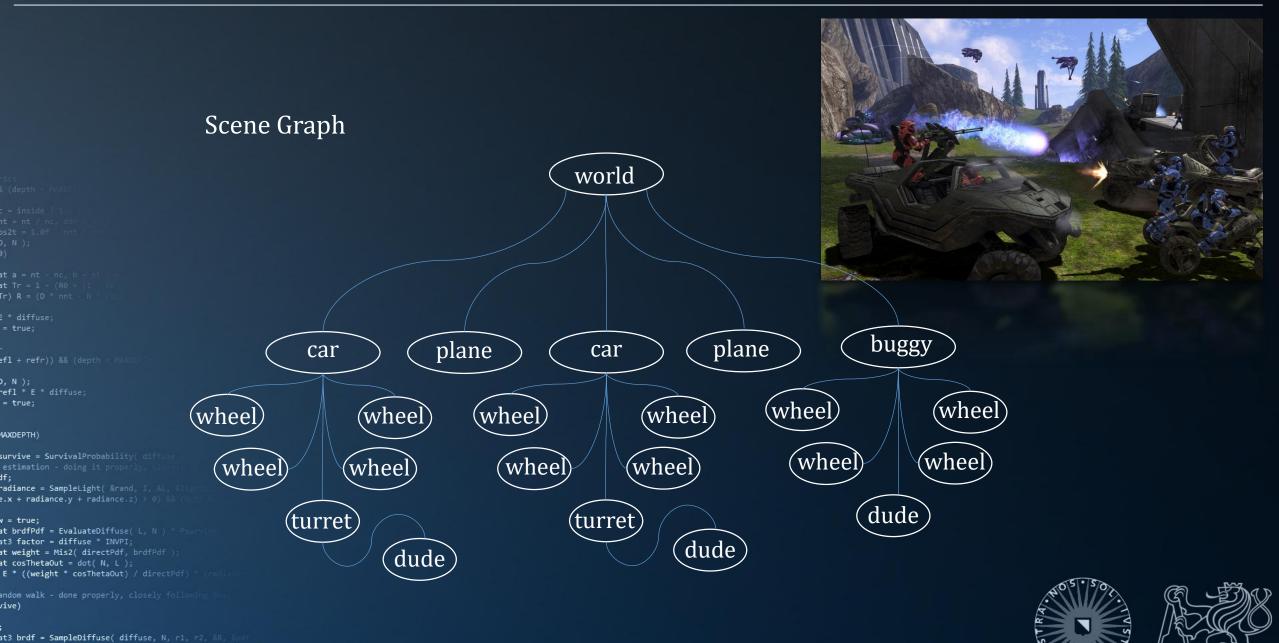
radiance = SampleLight(&rand, I, &L,

at brdfPdf = EvaluateDiffuse(L, N) * at3 factor = diffuse * INVPI;





pdf; n = E * brdf * (dot(N, R) / pdf);



Ray Tracing for Games

Scene Graph

If our application uses a scene graph, we can construct a BVH for each scene graph node.

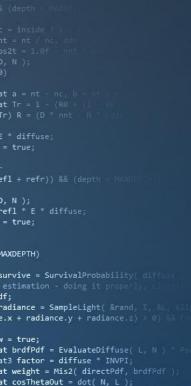
The BVH for each node is built using an appropriate construction algorithm:

- High-quality SBVH for static scenery (offline)
- Fast binned SAH BVHs for dynamic scenery

The extra nodes used to combine these BVHs into a single BVH are known as the *Top-level BVH*.







E * ((weight * cosThetaOut) / directPdf)
andom walk - done properly, closely follow

1 = E * brdf * (dot(N, R) / pdf);

at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, &

efl + refr)) && (depth <

survive = SurvivalProbability(diff

radiance = SampleLight(&rand, I, & e.x + radiance.y + radiance.z) > 0)

at brdfPdf = EvaluateDiffuse(L, N

refl * E * diffuse;

), N);

(AXDEPTH)

v = true;

The Top-level BVH - Construction

Input: *list of axis aligned bounding boxes for transformed scene graph nodes*

Algorithm:

- 1. Find the two elements in the list for which the AABB has the smallest surface area
- 2. Create a parent node for these elements
- 3. Replace the two elements in the list by the parent node
- 4. Repeat until one element remains in the list.

Note: algorithmic complexity is $O(N^3)$.



refl * E * diffuse;

survive = SurvivalProbability(diff

radiance = SampleLight(&rand, I, &

e.x + radiance.y + radiance.z) > 0

at brdfPdf = EvaluateDiffuse(L, N at3 factor = diffuse * INVPI; at weight = Mis2(directPdf, brdfPdf) at cosThetaOut = dot(N, L);

1 = E * brdf * (dot(N, R) / pdf);

at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, &

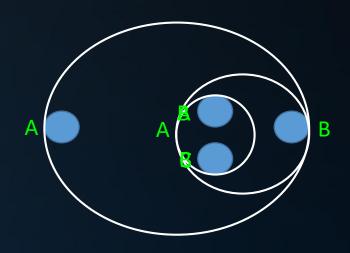
(AXDEPTH)

v = true;

The Top-level BVH – Faster Construction*

Algorithm:

```
Node A = list.GetFirst();
                  Node B = list.FindBestMatch( A );
                  while (list.size() > 1)
                      Node C = list.FindBestMatch( B );
                      if (A == C)
                         list.Remove( A );
                         list.Remove( B );
                         A = new Node(A, B);
                         list.Add( A );
                         B = list.FindBestMatch( A );
                      else A = B, B = C;
E * ((weight * cosThetaOut) / directPdf)
```







*: Fast Agglomerative Clustering for Rendering, Walter et al., 2008

- Combining BVHs
- Refitting
- Rigid Motion









Summary of BVH Characteristics

A BVH provides significant freedom.

- No need for a 1-to-1 relation between bounding boxes and primitives
- Bounding boxes may overlap
- Bounding boxes can be altered, as long as they fit in their parent box
- A BVH can be very bad but still valid

Some consequences / opportunities:

- We can rebuild part of a BVH
- We can combine two BVHs into one
- We can *refit* a BVH



```
E * ((weight * cosThetaOut) / directPdf) * (radian
andom walk - done properly, closely following Section
/ive)
;
st3 brdf = SampleDiffuse( diffuse, N, r1, r2, &R, &pourvive;
pdf;
n = E * brdf * (dot( N, R ) / pdf);
sion = true:
```

efl + refr)) && (depth

survive = SurvivalProbability(dif

radiance = SampleLight(&rand, I,

e.x + radiance.y + radiance.z) > 0)

at brdfPdf = EvaluateDiffuse(L, N)
at3 factor = diffuse * INVPI;
br weight = Mis2(directPdf, brdfPdf
at cosThetaOut = dot(N, L);

), N);

(AXDEPTH)

v = true;

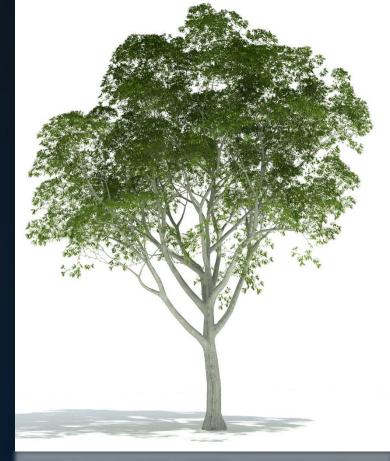
Refitting

Q: What happens to the BVH of a tree model, if we make it bend in the wind?

A: Likely, only bounds will change; the topology of the BVH will be the same (or at least similar) in each frame.

Refitting:

Updating the bounding boxes stored in a BVH to match changed primitive coordinates.





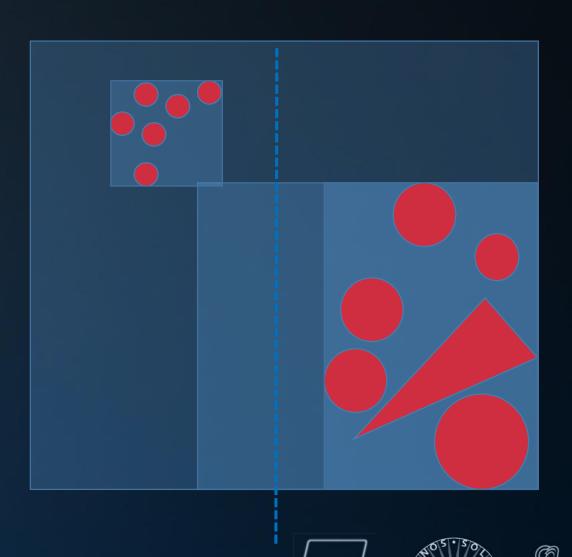


Refitting

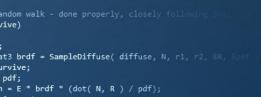
Updating the bounding boxes stored in a BVH to match changed primitive coordinates.

Algorithm:

- 1. For each leaf, calculate the bounds over the primitives it represents
- 2. Update parent bounds







(AXDEPTH)

v = true;

survive = SurvivalProbability(diff.

radiance = SampleLight(&rand, I, &L, e.x + radiance.y + radiance.z) > 0) &

at brdfPdf = EvaluateDiffuse(L, N) * at3 factor = diffuse * INVPI; at weight = Mis2(directPdf, brdfPdf); at cosThetaOut = dot(N, L);

E * ((weight * cosThetaOut) / directPdf)

Refitting - Suitability

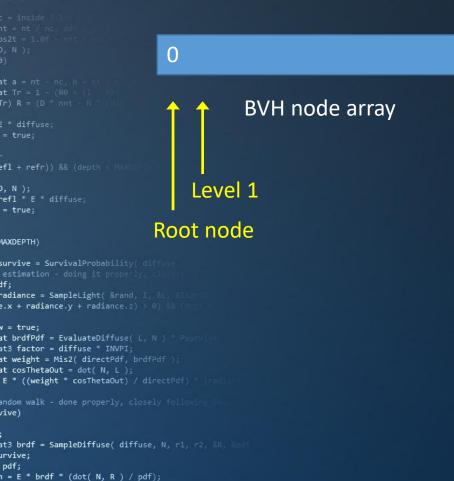








Refitting – Practical



Order of nodes in the node array:

We will never find the parent of node X at a position greater than X.

Therefore:

```
for( int i = N-1; i >= 0; i-- )
  nodeArray[i].AdjustBounds();
```





N-1

- Combining BVHs
- Refitting
- Rigid Motion





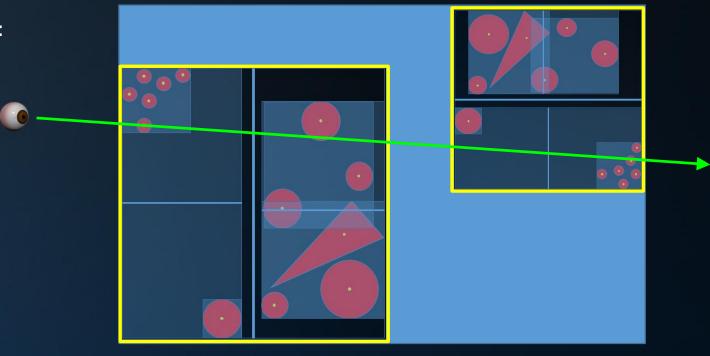


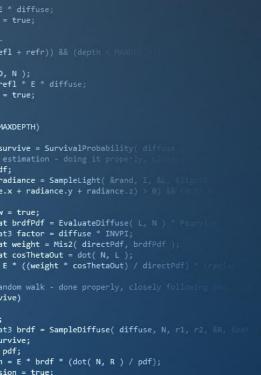
(AXDEPTH) survive = SurvivalProbability(diff) radiance = SampleLight(&rand, I, &L, &I e.x + radiance.y + radiance.z) > 0) 88 v = true; at brdfPdf = EvaluateDiffuse(L, N) Ps at3 factor = diffuse * INVPI; at weight = Mis2(directPdf, brdfPdf); at cosThetaOut = dot(N, L); at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, &pd ırvive; pdf; n = E * brdf * (dot(N, R) / pdf);

Rigid Motion

Applying rigid motion to a BVH:

- 1. Refit the top-level BVH
- 2. Refit the affected BVH







Rigid Motion

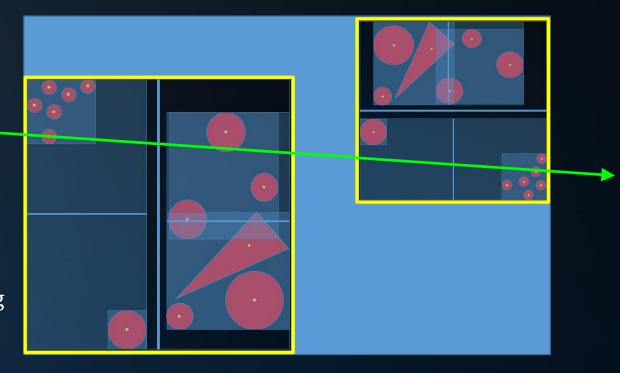
Applying rigid motion to a BVH:

- 1. Refit the top-level BVH
- 2. Refit the affected BVH

or:

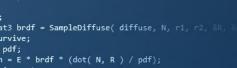
2. Transform the ray, not the node

Rigid motion is achieved by transforming the rays by the *inverse transform* upon entering the sub-BVH.



(this obviously does not only apply to translation)





andom walk - done properly, closely fol

), N);

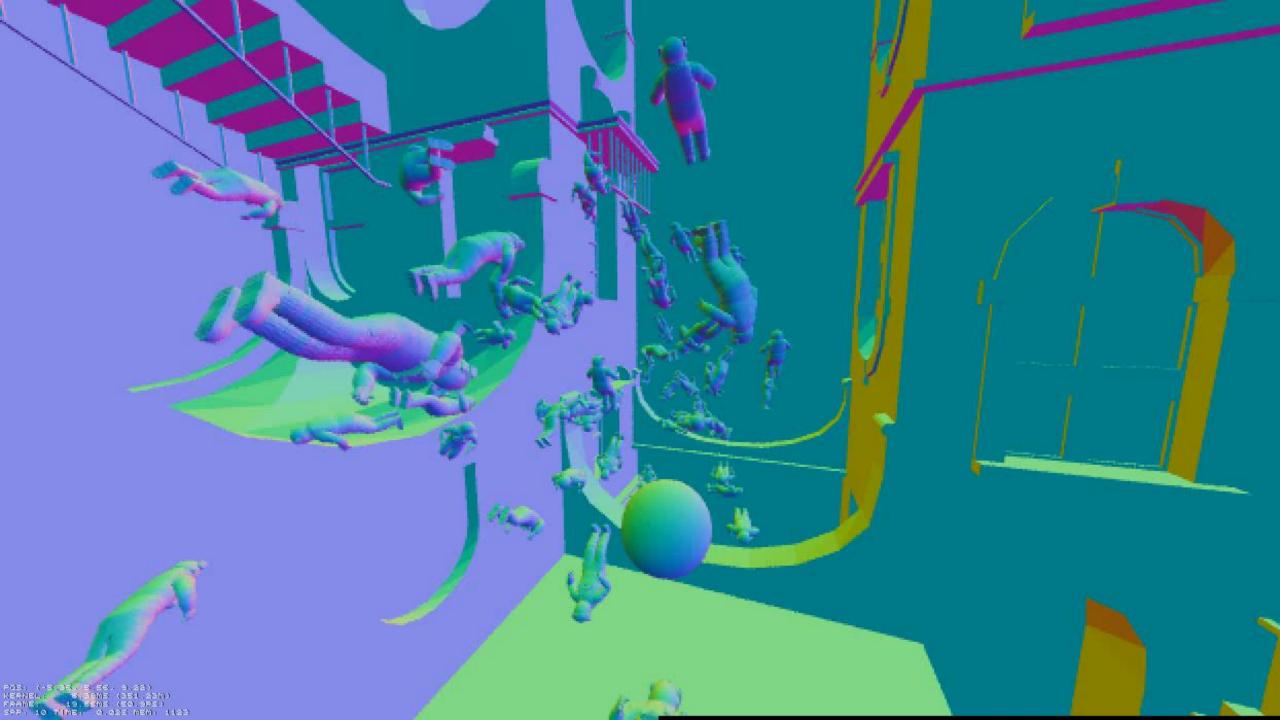
(AXDEPTH)

survive = SurvivalProbability(dif

radiance = SampleLight(&rand, I

at3 factor = diffuse * INVPI; at weight = Mis2(directPdf, brdfPdf at cosThetaOut = dot(N, L);

e.x + radiance.y + radiance.z) > 0



- Combining BVHs
- Refitting
- Rigid Motion







(AXDEPTH) survive = SurvivalProbability(diff) radiance = SampleLight(&rand, I, &L, &I e.x + radiance.y + radiance.z) > 0) 88 v = true; at brdfPdf = EvaluateDiffuse(L, N) Ps at3 factor = diffuse * INVPI; at weight = Mis2(directPdf, brdfPdf); at cosThetaOut = dot(N, L); at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, &pd ırvive; pdf; n = E * brdf * (dot(N, R) / pdf);

End of PART 6.



at a = nt - nc, b

(AXDEPTH)

v = true;

ırvive;

survive = SurvivalProbability(diffu

radiance = SampleLight(&rand, I, &L, &l 2.x + radiance.y + radiance.z) > 0) &&

at brdfPdf = EvaluateDiffuse(L, N) * P

1 = E * brdf * (dot(N, R) / pdf);

E * ((weight * cosThetaOut) / directPdf) * (re

at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, Apd

at3 factor = diffuse * INVPI; at weight = Mis2(directPdf, brdfPdf); at cosThetaOut = dot(N, L);





