```
at a = nt - nc, b
efl + refr)) && (depth < MAX
(AXDEPTH)
survive = SurvivalProbability( diff.
radiance = SampleLight( &rand, I, &L,
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;
1 = E * brdf * (dot( N, R ) / pdf);
```

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

Welcome!



Thursday 09:00 – 14:00

advanced Whitted audio, AI & physics faster Whitted Heaven7

LAB 2

work @ home

End result day 2:

A solid Whitted-style ray tracer, as a basis for subsequent work.

Friday 09:00 – 17:00

optimization profiling, rules of engagement threading



LAB 3

SIMD applied SIMD SIMD triangle SIMD AABB

LAB 4

YOU ARE HERE

Monday 09:00 – 17:00

acceleration grid, BVH, kD-tree SAH binning



LAB 5

GAME

JAM

refitting top-level BVH threaded building

LAB 6

Tuesday 09:00 – 17:00

Monte-Carlo Cook-style glossy, AA area lights, DOF



LAB 7

path tracing



LAB8

End result day 5:

Thursday 09:00 – 17:00

random numbers stratification blue noise



LAB 11

Friday

future work

09:00 - 17:00

importance path guiding sampling next event



LAB 10



End result day 3:

A 5x faster tracer.

End result day 4:

A real-time tracer.

Cook or Kajiya.

End result day 6:

Efficiency.

LAB9

estimation

LAB 10

End result day 6:

Great product.

Wednesday 13:00 – 17:00

efl + L D, N refl t

MAXDE survi esti Hf; radia

v = t at br at3 f at we at co E *

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

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

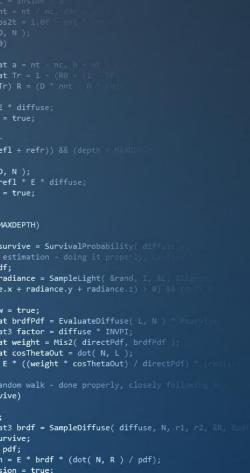
Agenda:

- Accelerate
- BVH
- Surface Area Heuristic
- Binning
- Fast Traversal







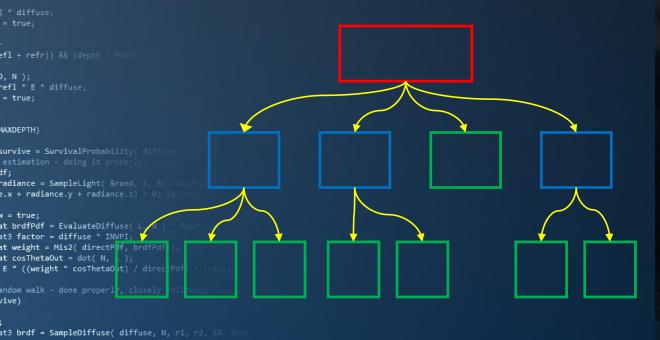


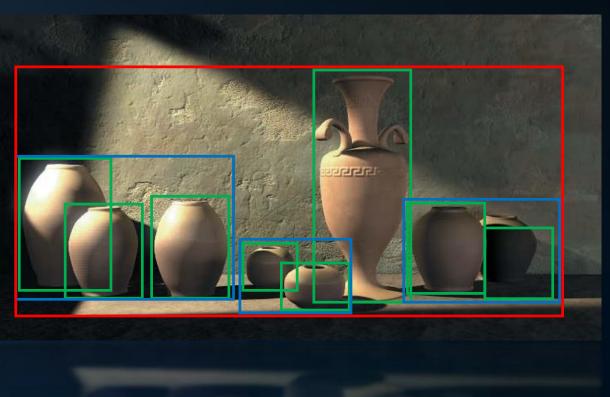
1 = E * brdf * (dot(N, R) / pdf);

The Bounding Volume Hierarchy

BVH: tree structure, with:

- a bounding box per node
- pointers to child nodes
- geometry at the leaf nodes





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

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

Automatic Construction of Bounding Volume Hierarchies

BVH: tree structure, with:

- a bounding box per node
- pointers to child nodes
- geometry at the leaf nodes

```
= true;

cef1 + refr)) && (depth < MAXDEFIN

D, N );

ref1 * E * diffuse;

= true;

MAXDEPTH)

Survive = SurvivalProbability( diffuse;
estination - doing it properly, closes and

if;
radiance = SampleLight( &rand, I, &E, &light
e.x + radiance.y + radiance.z) > 0) && (dot n

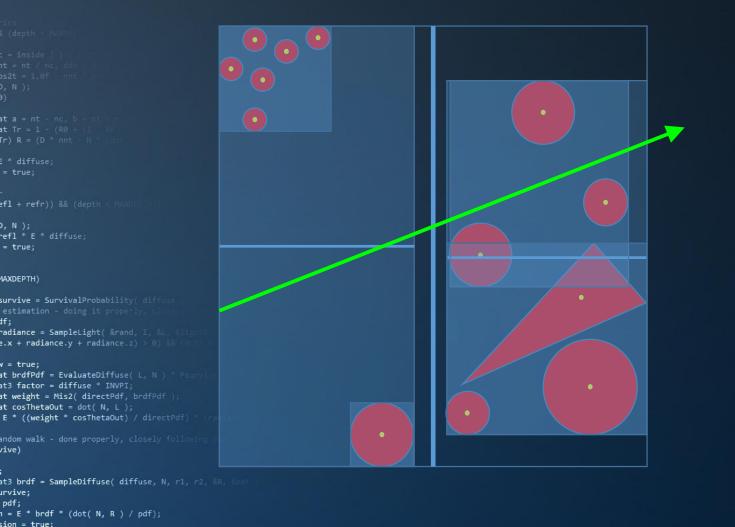
v = true;

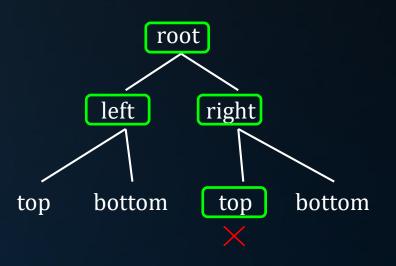
at brdfdf = EvaluateDiffuse( L, A ) * Paurvis

at weight = Mis2( directPlf, brdfPdf);
at cosThetaOut = dot( N, .);

E * ((Weight * cosThetaOut) / directPdf);* (radial sindow walk - done properly, closely following Ses
```

```
struct BVHNode
{
    AABB bounds;
    bool isLeaf;
    BVHNode*[] child;
    Primitive*[] primitive;
};
```





Automatic Construction of Bounding Volume Hierarchies

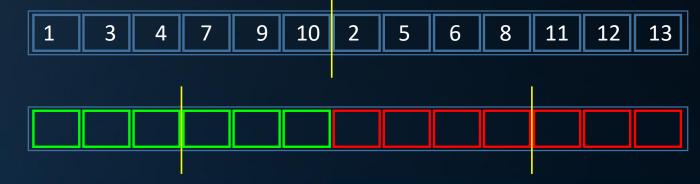


- 1. Determine AABB for primitives in array
- 2. Determine split axis and position
- 3. Partition
- 4. Repeat steps 1-3 for each partition

Note:

Step 3 can be done 'in place'.

This process is identical to QuickSort: the split plane is The 'pivot'.



/ive)
;
st3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, &pdf
urvive;
pdf;
n = E * brdf * (dot(N, R) / pdf);

at a = nt - nc,

), N);

(AXDEPTH)

v = true;

efl + refr)) && (depth

survive = SurvivalProbability(diff

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

at brdfPdf = EvaluateDiffuse(L, N)

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

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

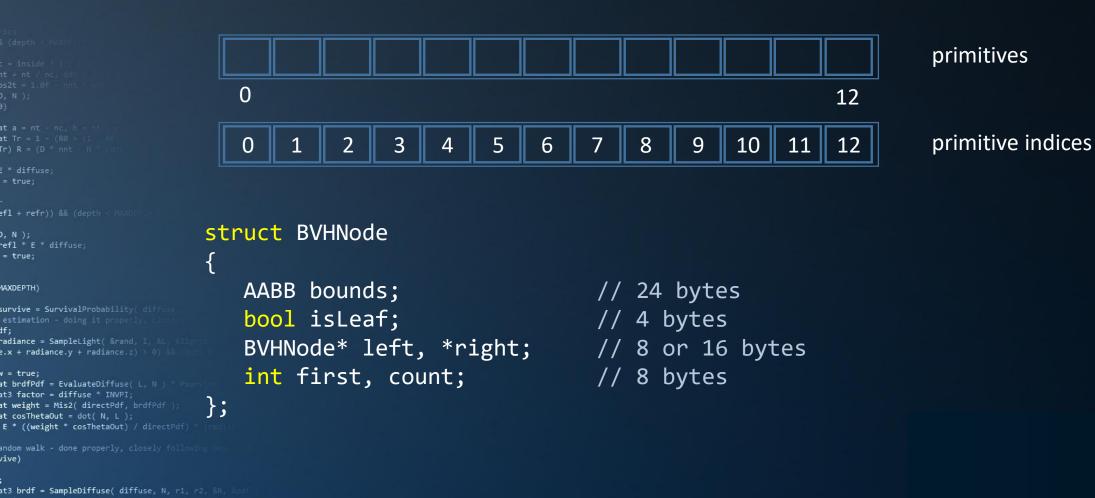
refl * E * diffuse;

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

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

```
12
                            struct BVHNode
                                 AABB bounds;
                                                                                   // 24 bytes
                                  bool isLeaf;
                                                                                   // 4 bytes
                                  BVHNode* left, *right; // 8 or 16 bytes
(AXDEPTH)
survive = SurvivalProbability( diff)
                                  Primitive** primList;  // ? bytes
radiance = SampleLight( &rand, I, &L
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
```

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

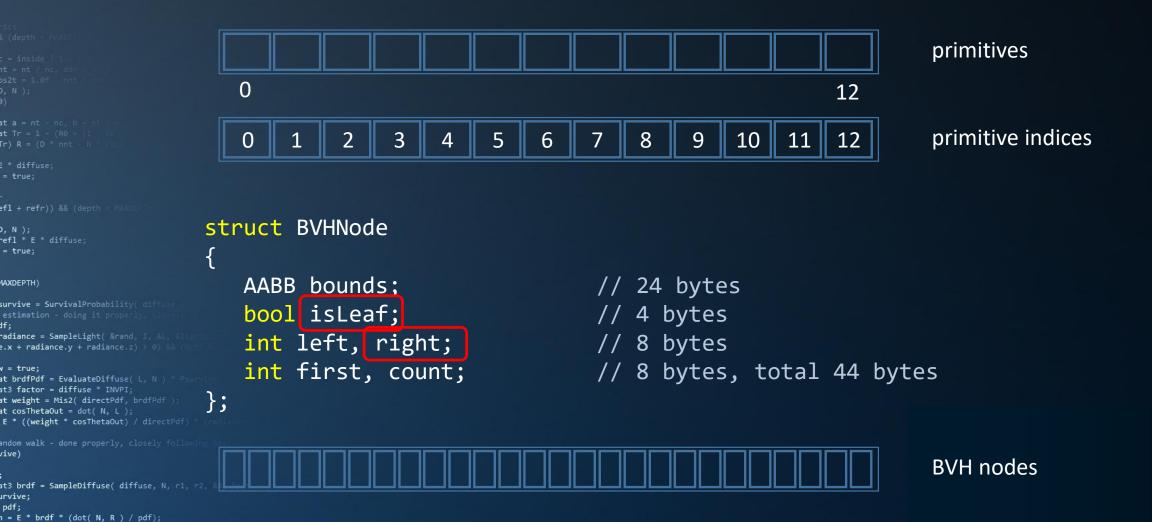


at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, &s

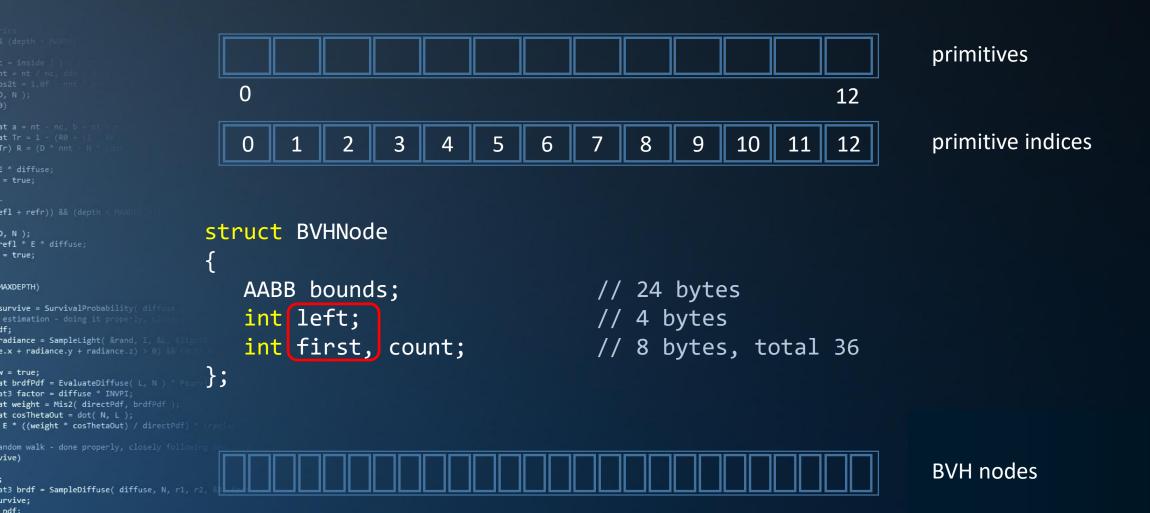
n = E * brdf * (dot(N, R) / pdf);

```
void BVH::ConstructBVH( Primitive* primitives )
                                                                                                        void BVHNode::Subdivide()
                                                                                                           if (count < 3) return;</pre>
                            // create index array
at a = nt - nc,
                            indices = new uint[N];
                                                                                                            this.left = new BVHNode();
                            for( int i = 0; i < N; i++ ) indices[i] = i;</pre>
                                                                                                            this.right = new BVHNode();
                                                                                                            Partition();
efl + refr)) && (depth <
                            // allocate BVH root node
                                                                                                            this.left->Subdivide();
), N );
                            root = new BVHNode();
                                                                                                            this.right->Subdivide();
refl * E * diffuse;
                                                                                                            this.isLeaf = false;
(AXDEPTH)
                            // subdivide root node
survive = SurvivalProbability( diff
                            root->first = 0;
radiance = SampleLight( &rand, I, &
                            root->count = N;
e.x + radiance.y + radiance.z) > 0)
                            root->bounds = CalculateBounds( primitives, root->first, root->count );
v = true;
at brdfPdf = EvaluateDiffuse( L.
at3 factor = diffuse * INVPI;
                            root->Subdivide();
at weight = Mis2( directPdf, brdfPdf
at cosThetaOut = dot( N, L );
E * ((weight * cosThetaOut) / directPdf)
andom walk - done properly, closely follow
```

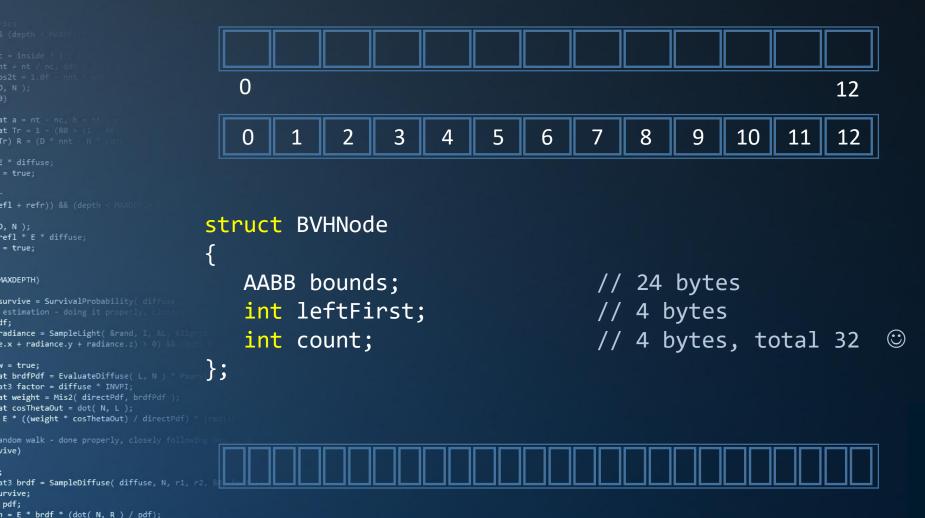
```
void BVH::ConstructBVH( Primitive* primitives )
                                                                                                          void BVHNode::Subdivide()
                                                                                                               if (count < 3) return;</pre>
                             // create index array
at a = nt - nc,
                             indices = new uint[N];
                                                                                                               this.left = &pool[poolPtr++];
                                                                                                               this.right = &pool[poolPtr++];
                             for( int i = 0; i < N; i++ ) indices[i] = i;</pre>
                                                                                                               Partition();
efl + refr)) && (depth < )
                             // allocate BVH root node
                                                                                                               this.left->Subdivide();
), N );
                             pool = new BVHNode[N * 2];
                                                                                                               this.right->Subdivide();
refl * E * diffuse;
                             root = &pool[0];
                                                                                                               this.isLeaf = false;
(AXDEPTH)
                             poolPtr = 2;
survive = SurvivalProbability( diff
radiance = SampleLight( &rand, I, &L
                             // subdivide root node
e.x + radiance.y + radiance.z) > 0)
                             root->first = 0;
v = true;
at brdfPdf = EvaluateDiffuse( L, N
                             root->count = N;
at3 factor = diffuse * INVPI;
at weight = Mis2( directPdf, brdfPdf )
at cosThetaOut = dot( N, L );
                             root->bounds = CalculateBounds( primitives, root->first, root->count );
E * ((weight * cosThetaOut) / directPdf)
                             root->Subdivide();
andom walk - done properly, closely follo
at3 brdf = SampleDiffuse( diffuse, N, r1, r2, 8
n = E * brdf * (dot( N, R ) / pdf);
```



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

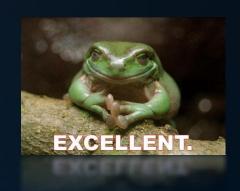


Automatic Construction of Bounding Volume Hierarchies



primitives

primitive indices



BVH nodes

TL;DR

Building a BVH:

- We build a BVH over triangle indices.
- The optimal BVH node structure is 32 bytes in size.
- Node 'pointers' are actually indices in a pre-allocated, aligned array.
- Node 1 is unused.

```
struct BVHNode
{
    float3 bmin;
    int leftFirst;
    float3 bmax;
    int count;
};
```

```
AXXDEPTH)

survive = SurvivalProbability( diffuse estimation - doing it properly, close)

If;

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

v = true;

at brdfPdf = EvaluateDiffuse( L, N ) * Psust3 factor = diffuse * INVPI;

at weight = Mis2( directPdf, brdfPdf );

at weight = Mis2( directPdf, brdfPdf );

E * ((weight * cosThetaOut) / directPdf)

sandom walk - done properly, closely followive)

int3 brdf = SampleDiffuse( diffuse, N, rl, restrictions)
```

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

```
AL MINISTER PRINTS

TO THE PRINTS

T
```

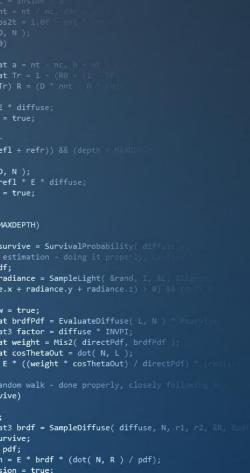
Agenda:

- Accelerate
- BVH
- Surface Area Heuristic
- Binning
- Fast Traversal

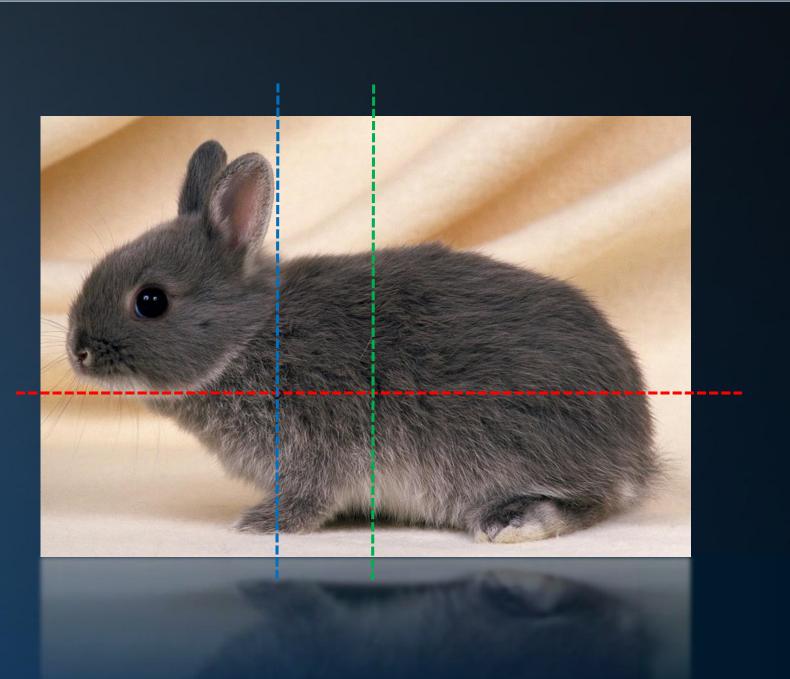




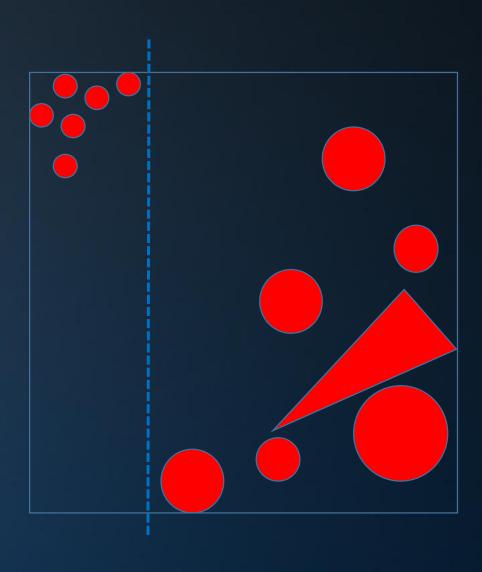




```
(AXDEPTH)
radiance = SampleLight( &rand, I, &L, &l
e.x + radiance.y + radiance.z) > 0) && (
v = true;
at brdfPdf = EvaluateDiffuse( L, N ) * Pst
at3 factor = diffuse * INVPI;
at weight = Mis2( directPdf, brdfPdf );
at cosThetaOut = dot( N, L );
E * ((weight * cosThetaOut) / directPdf) * (radd
/ive)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



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



What Are We Trying To Solve?

A BVH is used to reduce the number of ray/primitive intersections.

But: it introduces new intersections.

The ideal BVH minimizes:

- # of ray / primitive intersections
- # of ray / node intersections.

Optimal Split Plane Position

The ideal split minimizes the *expected cost* of a ray intersecting the resulting nodes.

This expected cost is based on:

- Number of primitives that will have to be intersected
- Probability of this happening

The cost of a split is thus:

$$A_{left} * N_{left} + A_{right} * N_{right}$$



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

Optimal Split Plane Position

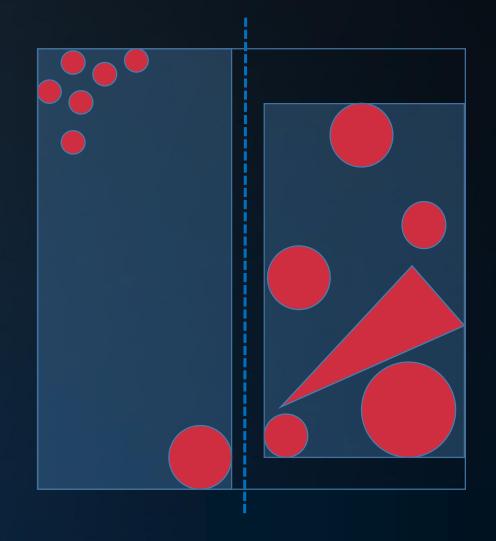
The ideal split minimizes the *expected cost* of a ray intersecting the resulting nodes.

This expected cost is based on:

- Number of primitives that will have to be intersected
- Probability of this happening

The cost of a split is thus:

$$A_{left} * N_{left} + A_{right} * N_{right}$$



```
radiance = SampleLight( &rand, I, &L, light)

e.x + radiance.y + radiance.z) > 0) && (decompose)

w = true;

at brdfPdf = EvaluateDiffuse( L, N ) * Psurvive

at3 factor = diffuse * INVPI;

at weight = Mis2( directPdf, brdfPdf );

at cosThetaOut = dot( N, L );

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

andom walk - done properly, closely following securive)

;

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

urvive;

pdf;

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

sion = true:
```

survive = SurvivalProbability(diff

(AXDEPTH)

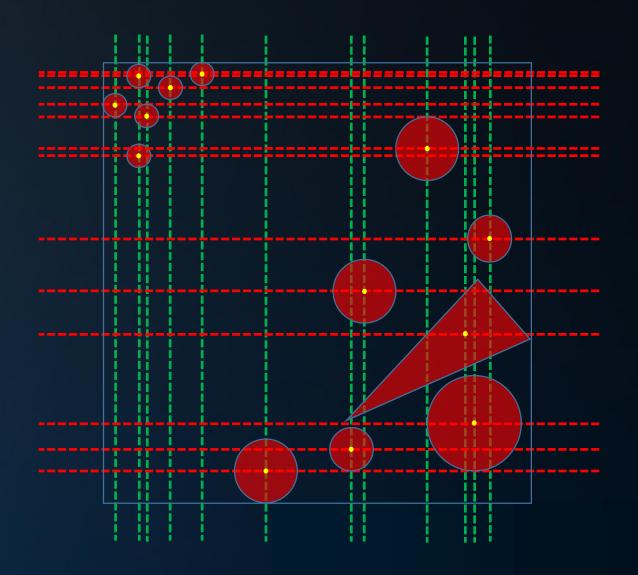
Optimal Split Plane Position

Which positions do we consider?

Object subdivision may happen over x, y or z axis.

The cost function is constant between primitive centroids.

→ For N primitives: 3(N-1) possible locations



e.x + radiance.y + radiance.z) > 0) && (detail
v = true;
at brdfPdf = EvaluateDiffuse(L, N) * Psurvive
at3 factor = diffuse * INVPI;
at weight = Mis2(directPdf, brdfPdf);
at cosThetaOut = dot(N, L);
E * ((weight * cosThetaOut) / directPdf) * (radian
andom walk - done properly, closely following servive)
;
at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, &p
urvive;
pdf;
n = E * brdf * (dot(N, R) / pdf);
sion = true;

(AXDEPTH)

survive = SurvivalProbability(dif

radiance = SampleLight(&rand, I, &L

(AXDEPTH)

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

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

A split is 'not worth it' if it doesn't yield

$$A_{left} * N_{left} + A_{right} * N_{right} \ge A * N$$

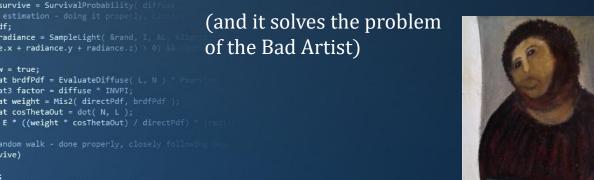
This provides us with a natural and optimal termination criterion.

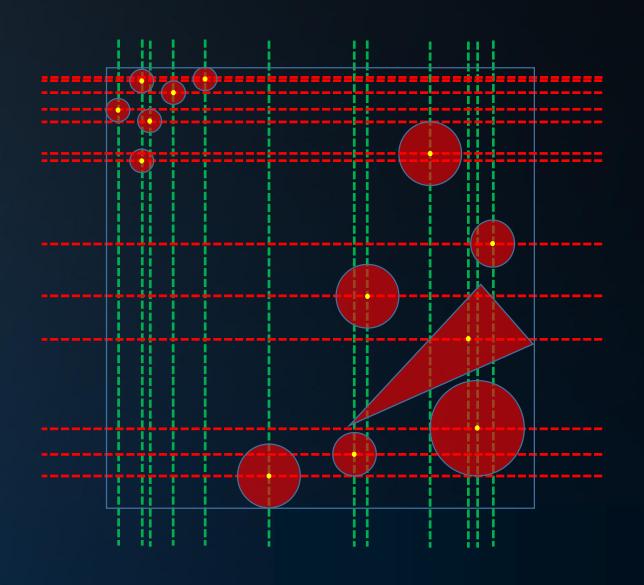




a cost lower than the cost of the parent node, i.e.:

$$A_{left} * N_{left} + A_{right} * N_{right} \ge A * N$$

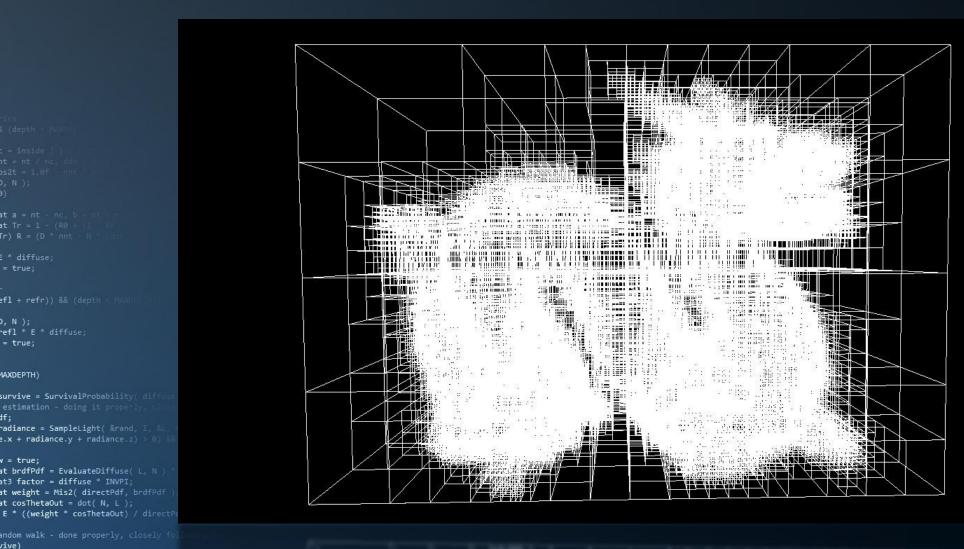




at3 brdf = SampleDiffuse(diffuse, N, r1, r2, 8

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

urvive; pdf;

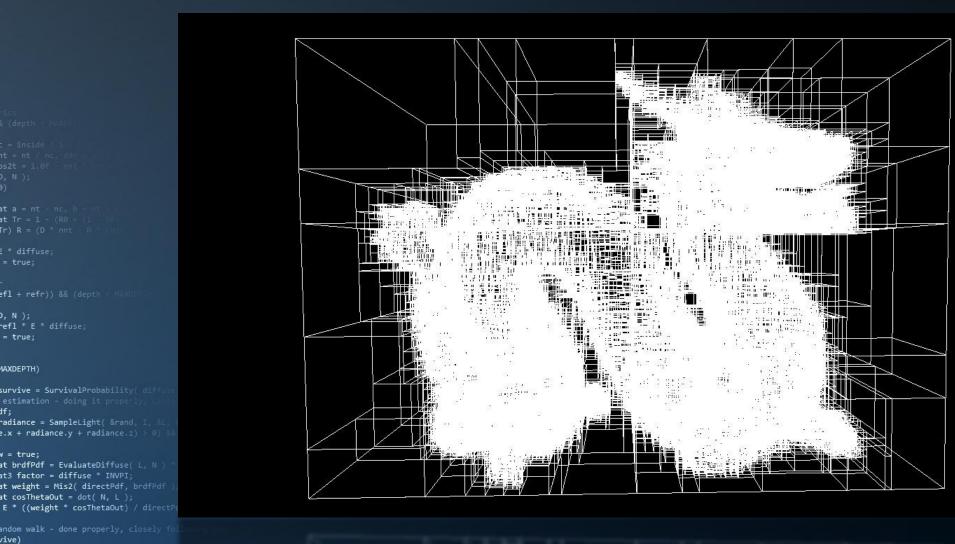


Median Split

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

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

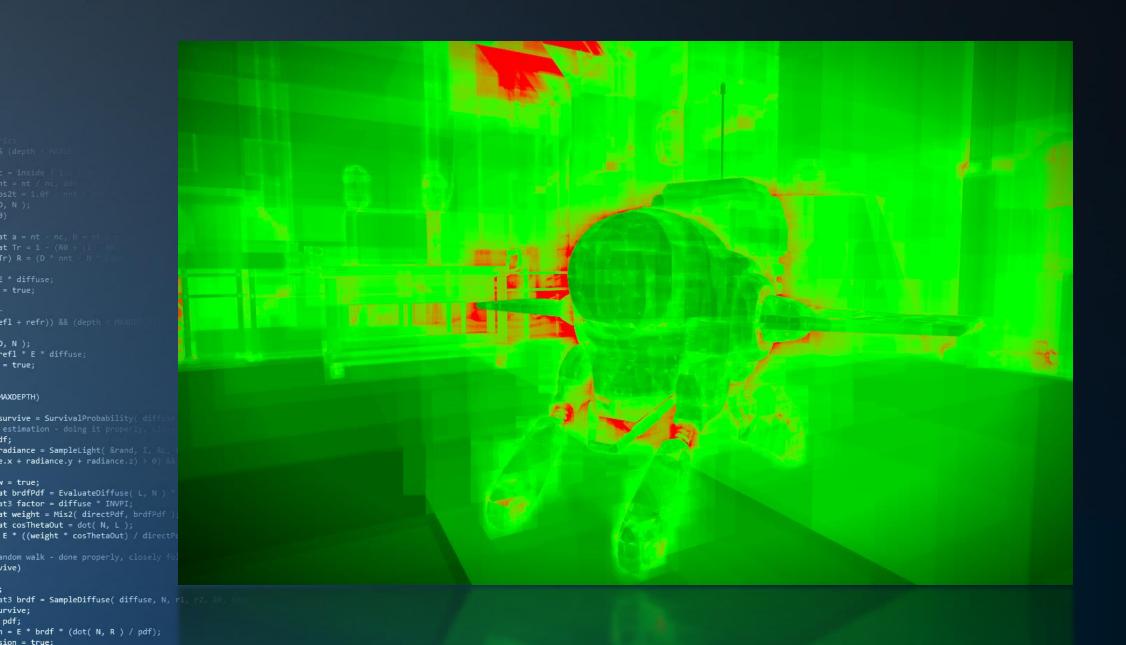
urvive; pdf;

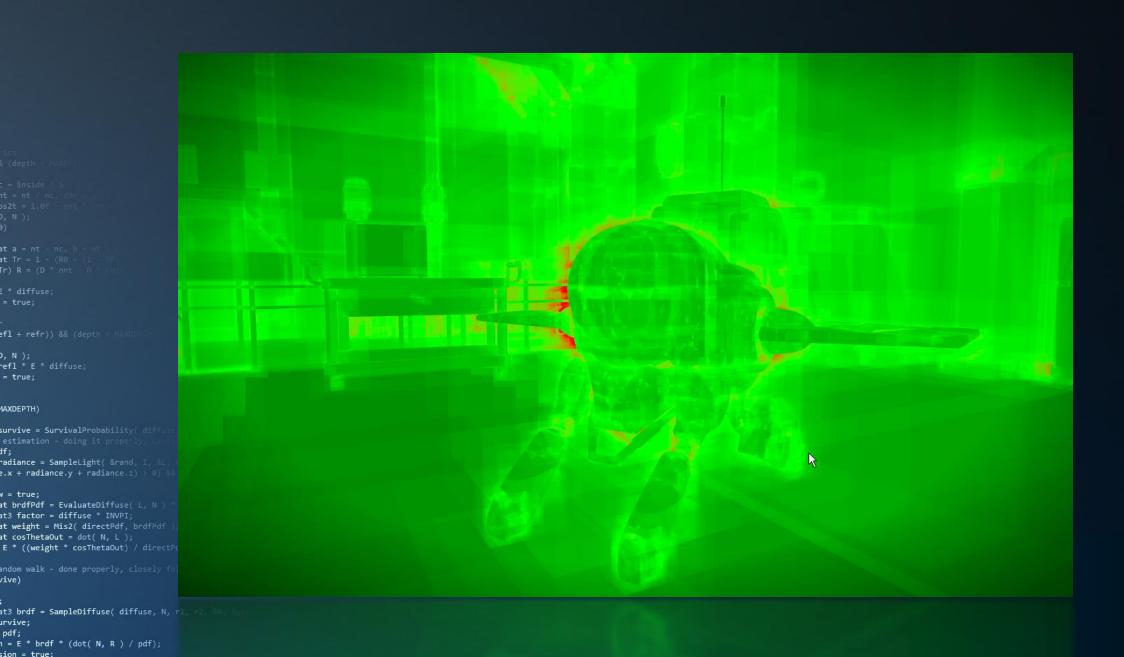


Surface Area Heuristic

```
(AXDEPTH)
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) / directF
```

at3 brdf = SampleDiffuse(diffuse, N, r1, r2, & urvive; pdf; n = E * brdf * (dot(N, R) / pdf);





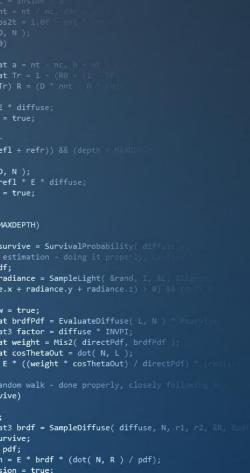
Agenda:

- Accelerate
- BVH
- Surface Area Heuristic
- Binning
- Fast Traversal









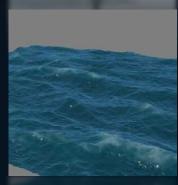
Rapid BVH Construction

Rebuilding a BVH requires 3*NlogN* split plane evaluations.

Speeding it up:

- 1. Do not use SAH (significantly lower quality BVH)
- 2. Do not evaluate all 3 axes (minor degradation of BVH quality)
- 3. Make split plane selection independent of *N*

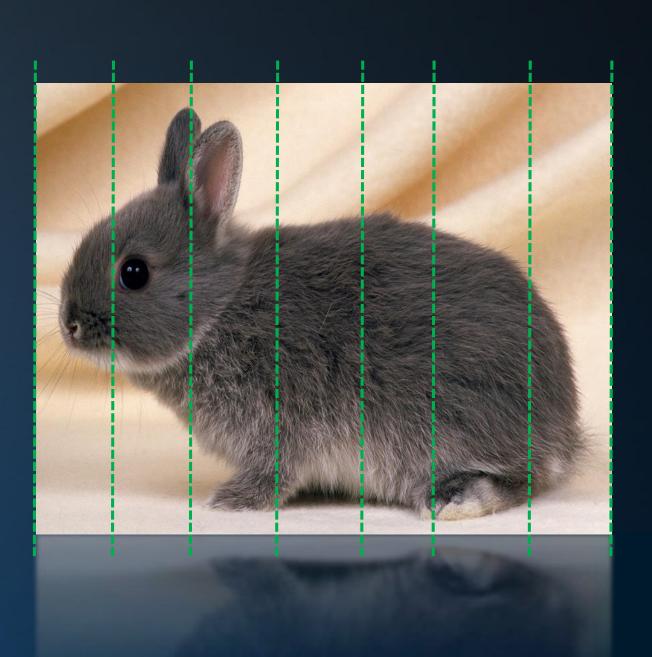






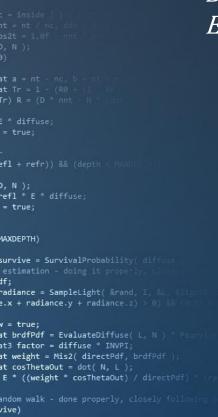


```
(AXDEPTH)
radiance = SampleLight( &rand, I, &L, &l)
e.x + radiance.y + radiance.z) > 0) && (
v = true;
at brdfPdf = EvaluateDiffuse( L, N ) * Psu
at3 factor = diffuse * INVPI;
at weight = Mis2( directPdf, brdfPdf );
at cosThetaOut = dot( N, L );
E * ((weight * cosThetaOut) / directPdf) * (radd
/ive)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



Binned BVH Construction*

Binned construction: Evaluate SAH at N discrete intervals.





*: On fast Construction of SAH-based Bounding Volume Hierarchies, Wald, 2007

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

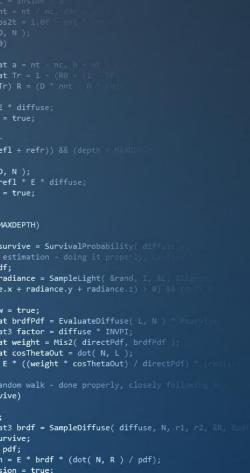
Agenda:

- Accelerate
- BVH
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- Fast Traversal









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

BVH Traversal

```
Ray:
                              Basic process:
                                                                                       vec3 O, D
                              BVHNode::Traverse( Ray r )
                                                                                       float t
                                   if (!r.Intersects( bounds )) return;
                                   if (isleaf())
                                        IntersectPrimitives();
                                   else
(AXDEPTH)
survive = SurvivalProbability( diff)
                                        pool[left].Traverse( r );
radiance = SampleLight( &rand, I, &L
                                        pool[left + 1].Traverse( r );
e.x + radiance.y + radiance.z) > 0) 8
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 follow
at3 brdf = SampleDiffuse( diffuse, N, r1, r2, &R, Upd
```

Ordered traversal, option 1:

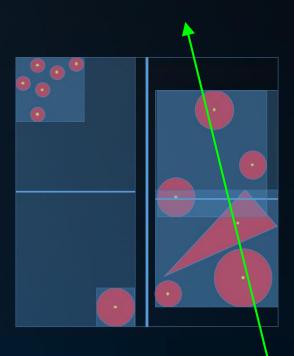
- Calculate distance to both child nodes
- Traverse the nearest child node first

Ordered traversal, option 2:

- For each BVH node, store the axis along which it was split
- Use ray direction sign for that axis to determine near and far

Ordered traversal, option 3:

- Determine the axis for which the child node centroids are furthest apart
- Use ray direction sign for that axis to determine near and far.



at3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, A 1 = E * brdf * (dot(N, R) / pdf);

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;

at weight = Mis2(directPdf, brdfPdf at cosThetaOut = dot(N, L);

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

refl * E * diffuse;

), N);

(AXDEPTH)

v = true;

BVH Traversal

AAXDEP

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

Ray/AABB Intersection

```
Vector code:
bool intersection( box b, ray r )
{
    __m128 t1 = _mm_mul_ps( _mm_sub_ps( node->bmin4, O4 ), rD4 );
    __m128 t2 = _mm_mul_ps( _mm_sub_ps( node->bmax4, O4 ), rD4 );
    __m128 vmax4 = _mm_max_ps( t1, t2 ), vmin4 = _mm_min_ps( t1, t2 )
    float* vmax = (float*)&vmax4, *vmin = (float*)&vmin4;
    float tmax = min(vmax[0], min(vmax[1], vmax[2]));
    float tmin = max(vmin[0], max(vmin[1], vmin[2]));
    return tmax >= tmin && tmax >= 0;
}

struct BVHNode
{
```

AABB bounds;

int count;

int leftFirst;

```
struct BVHNode
{
    float3 bmin;
    int leftFirst;
    float3 bmax;
    int count;
};
```

```
struct BVHNode
   union
      struct
         float3 bmin;
         int leftFirst;
        m128 bmin4;
   union
      struct
         float3 bmax;
         int count;
      m128 bmax4;
```

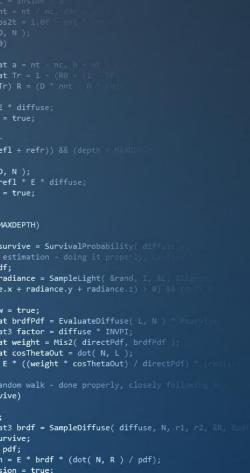
Agenda:

- Accelerate
- BVH
- Surface Area Heuristic
- Binning
- Fast Traversal



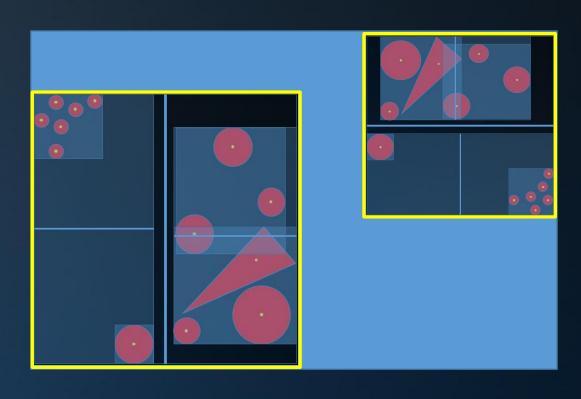






Combining BVHs

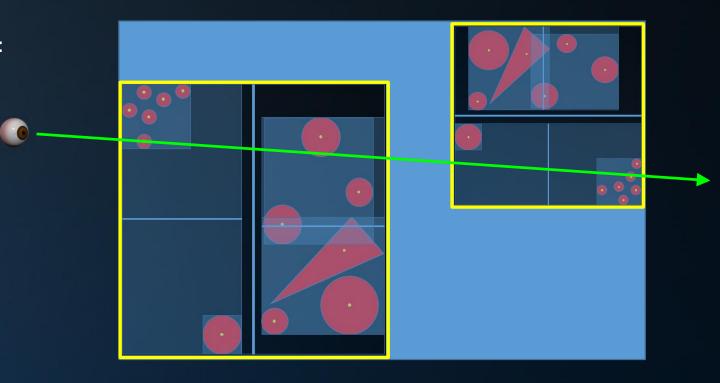
```
(AXDEPTH)
survive = SurvivalProbability( diffus
radiance = SampleLight( &rand, I, &L, &L
e.x + radiance.y + radiance.z) > 0) 88
v = true;
at brdfPdf = EvaluateDiffuse( L, N ) * Ps
at3 factor = diffuse * INVPI;
at weight = Mis2( directPdf, brdfPdf );
at cosThetaOut = dot( N, L );
E * ((weight * cosThetaOut)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



Rigid Motion

Applying rigid motion to a BVH:

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



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

Rigid Motion

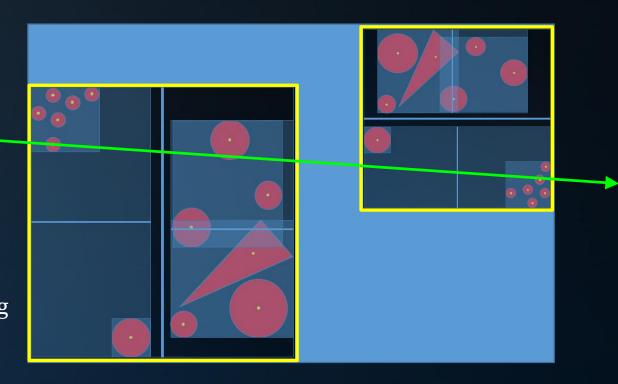
Applying rigid motion to a BVH:

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

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)

nt3 brdf = SampleDiffuse(diffuse, N, r1, r2, &R, & prvive; pdf; n = E * brdf * (dot(N, R) / pdf);

andom walk - done properly, closely fol

survive = SurvivalProbability(dif

radiance = SampleLight(&rand, I,

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

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

), N);

(AXDEPTH)

v = true;

End of PART 5.









