

Precomputed Dynamic Appearance Synthesis and Rendering

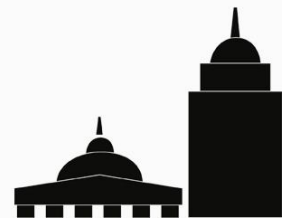
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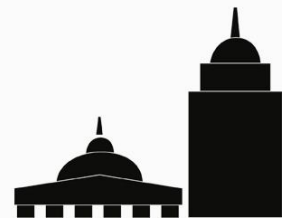
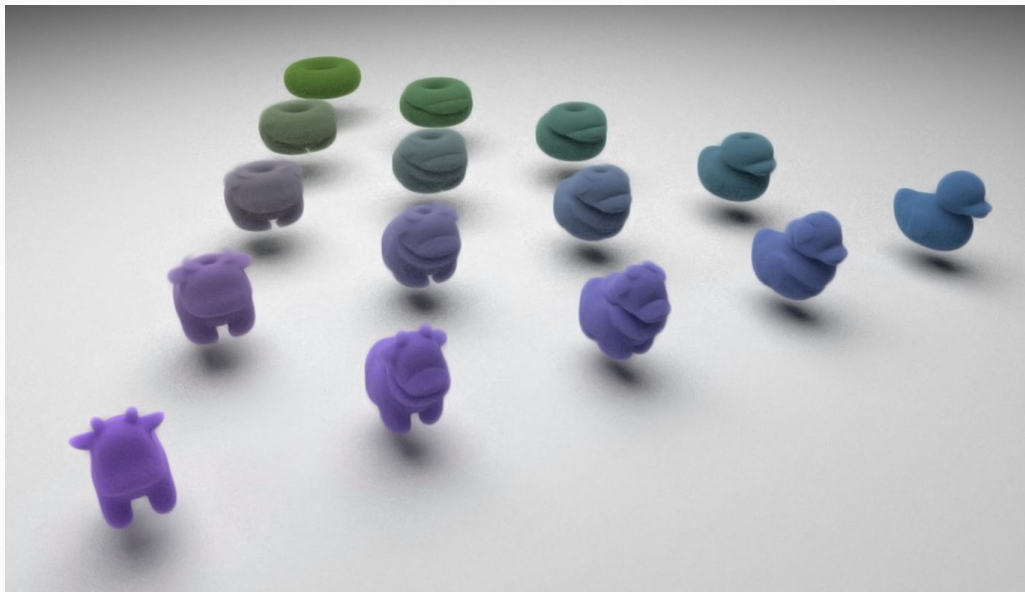
²Adobe Research, USA

Invited presenter: Iliyan Georgiev³

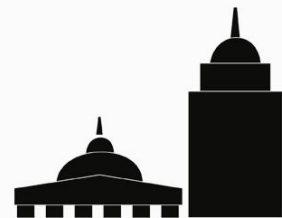
³Adobe Research, UK



Teasers



Background



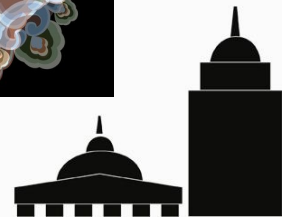
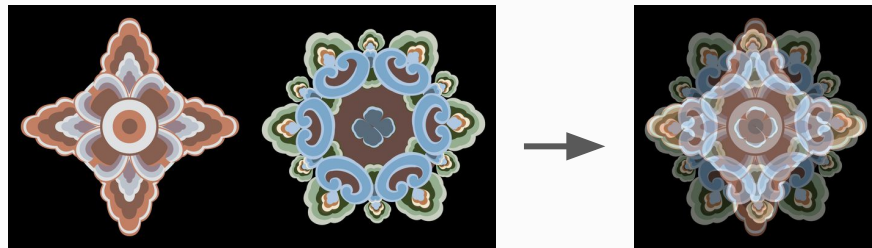
Appearances

Appearance in computer graphics: textures, volumes, BRDFs, etc.

Appearance synthesis: interpolating existing appearances to create additional materials.

Existing issues:

- Ghosting artifact;
- Efficiency;
- Storage.



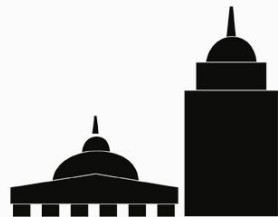
Motivation & Target

Our goal:

Integrating a **general**, efficient, and cross-dimensional **natural** appearance blending and rendering solution into traditional renderers.

Requirements:

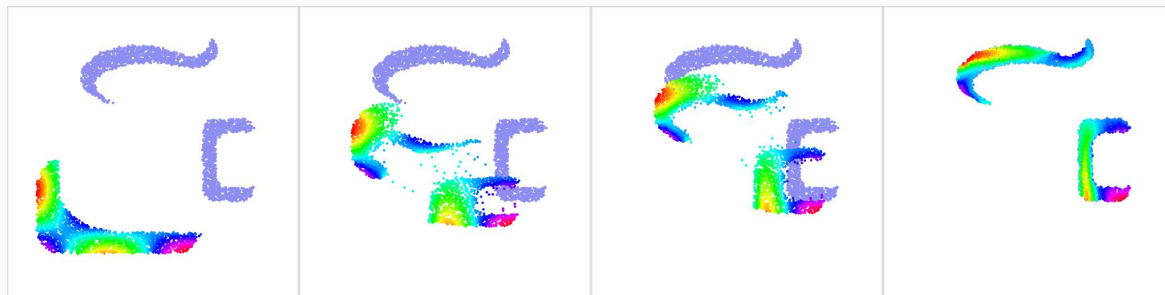
- Natural distribution interpolation;
- High performance;
- Point-sample and/or importance-sample capability;
- No full data reconstruction needed: local query;
- Varying interpolation weights.



Optimal Transport (OT)

Mathematical tool:

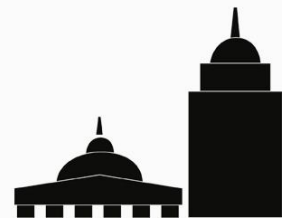
- Offers promising solutions for natural interpolation between distributions;
- Provide CLOSEST distances between distributions:
 - minimum sum of moving distance
 - total “weight” conservation
- Discrete or continuous;
- Dimension-agnostic.



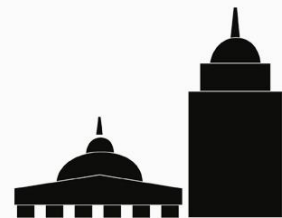
OT in Rendering

Existing OT solvers:

- Computationally intensive;
- No existing point query support for OT;
- Rendering integration:
 - Monte Carlo point-sampling structure
 - work with discrete point sets
 - one-to-one mapping



Our Method



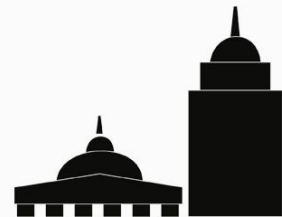
Our Method

Step 1: Precomputation:

- A common proxy distribution;
- Calculate OT between each input and the proxy.

Step 2: Runtime renderer integration:

- Construct hierarchical query structure;
- Point query or range query to quickly find mapping;
- Interpolation based on blending weight.



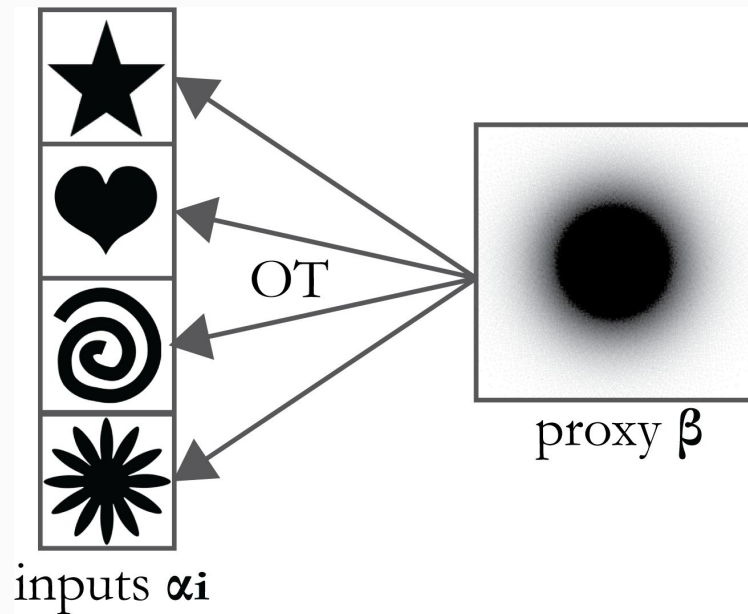
Step 1: Precomputation

Target:

- From common proxy distribution β to different inputs α_i
 - one-to-one discrete sample mapping

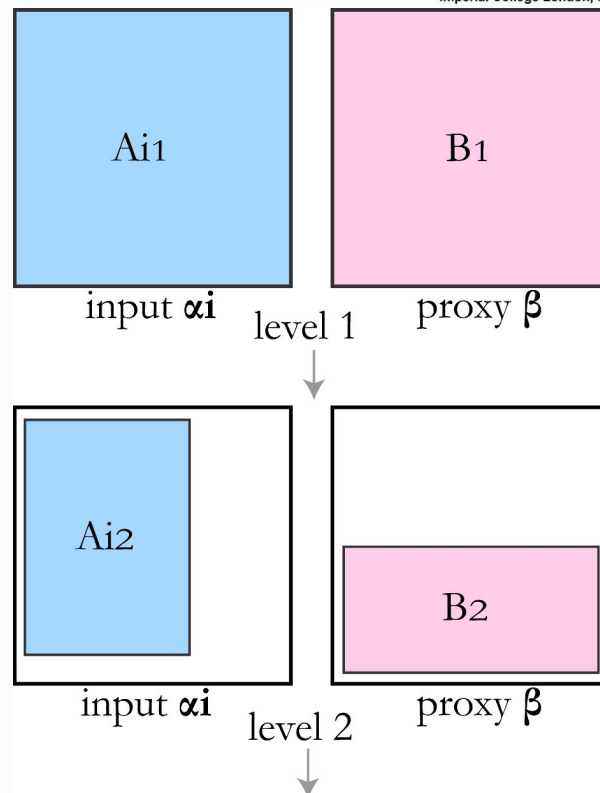
OT solver:

- Geomloss: by Feydy et al.;
- Runs on GPU;
- Discrete one-to-one point mapping.



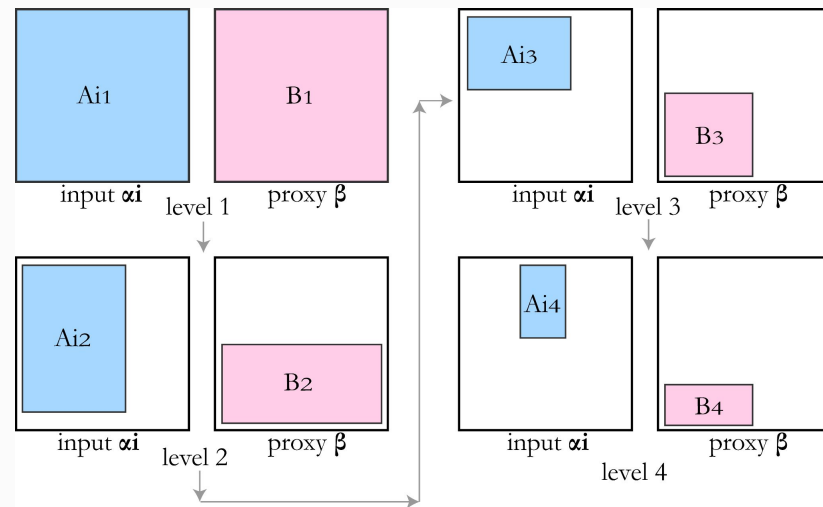
Step 2-1: Hierarchical Query Structure

- Inspired by bounding volume hierarchy (BVH);
- Each level contains:
 - two axis-aligned AABB bounding boxes for samples A_{ij} (j stands for the j -th hierarchical level)
 - all samples in α_i (i stands for the i -th input) and β inside the bounding box
 - two children pointers

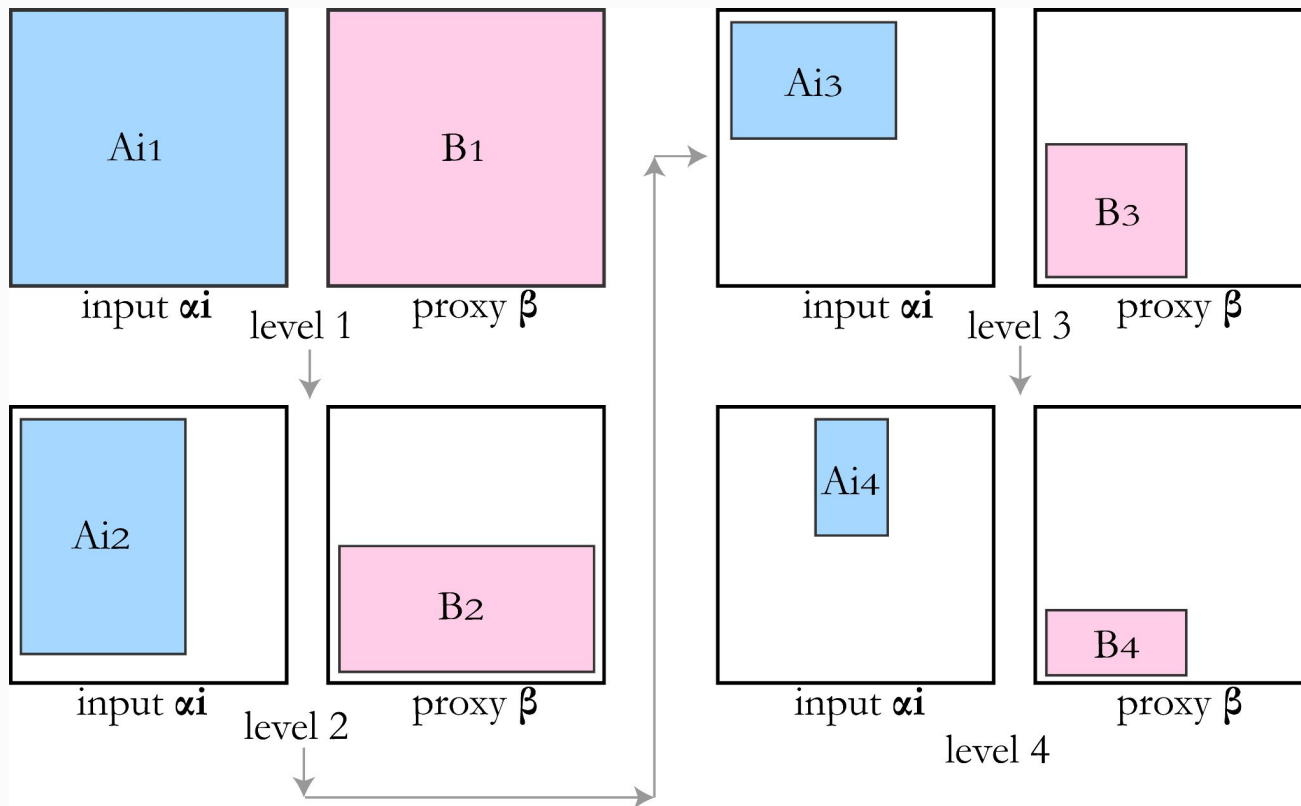


Step 2-1: Hierarchical Query Structure

- Level subdivision on β :
 - based on the longest Euclidean distance of samples of one dimension
 - divide the proxy bounding box B_j in half
 - find mapped samples from input α_i
 - calculate input bounding box A_{ij}
- Subdivision until only one sample;
- Facilitates range queries calculation.

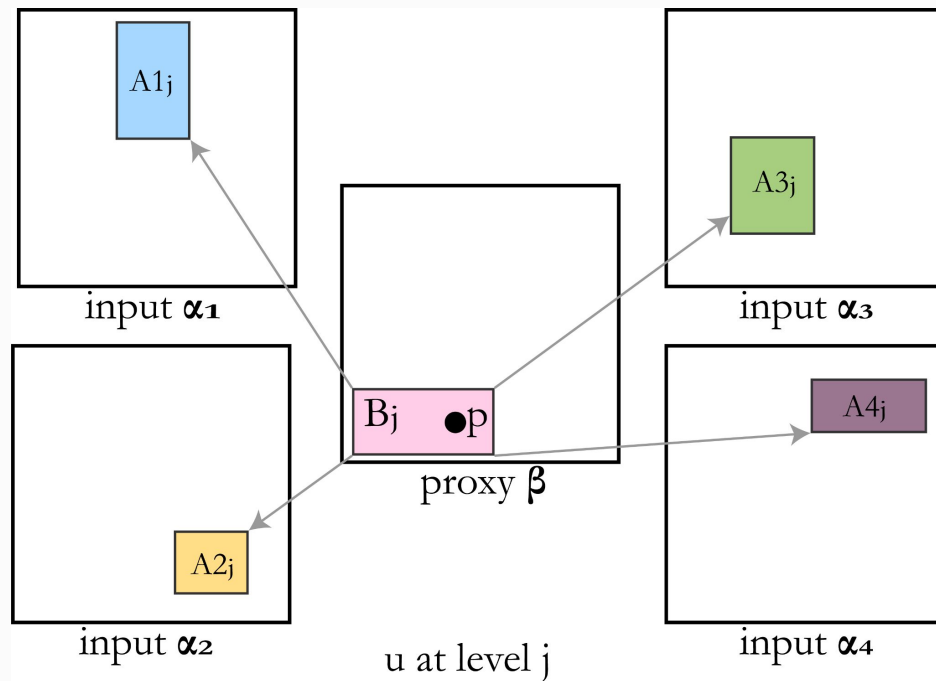


Step 2-1: Hierarchical Query Structure



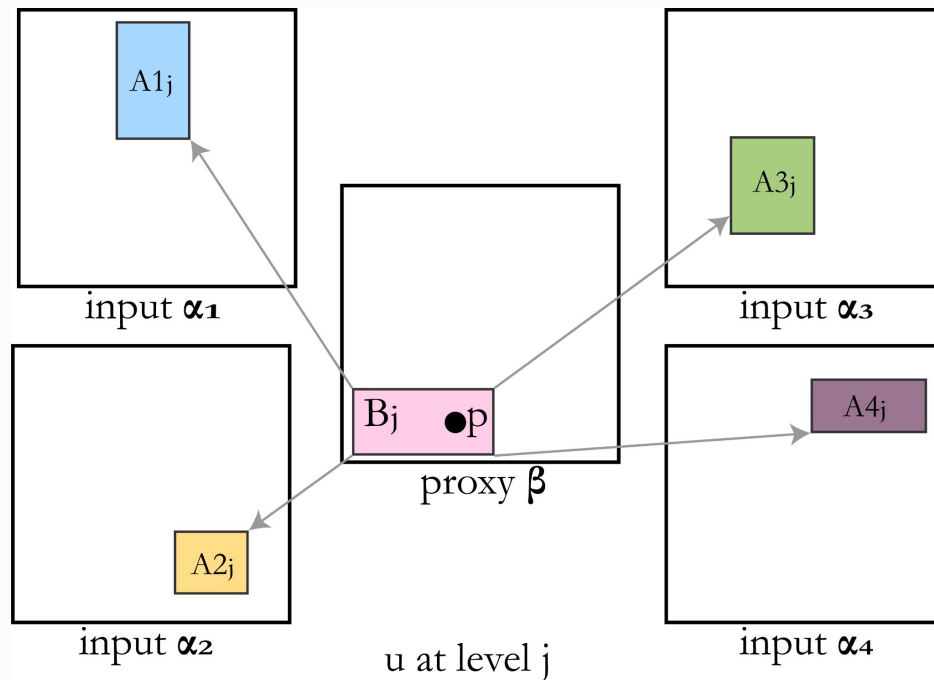
Step 2-2: Runtime Blending

- For a spatial point p that takes blending weight u :
 - traverse the hierarchical structure
 - find mapped bounding box at level j
 - point query: find the leaf nodes and perform blending
 - range query: find the largest covered bounding box and perform blending



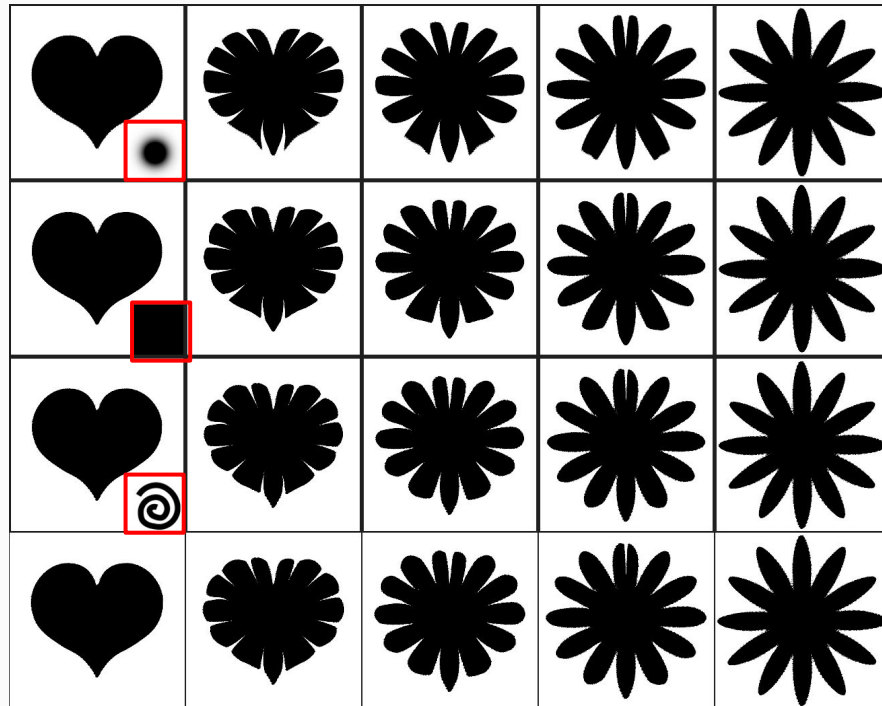
Step 2-2: Runtime Blending

- No limitation on amount of inputs given proper u ;
- Flexibility to switch inputs;
- Query allows data generation only around the spatial point p .



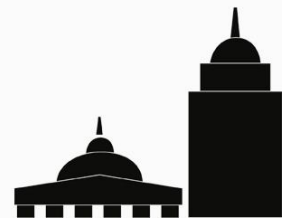
Proxy Selection

- Different proxies:
 - Gaussian
 - Uniform
 - Spiral
 - No proxy
- The differences in using different proxies or without proxy are minimal.

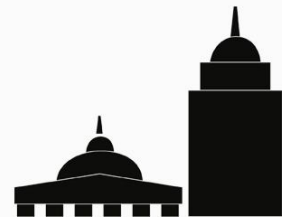


Implementation

- Geomloss OT solver by Feydy et al.:
 - on GPU
 - precomputation only
- Mitsuba renderer:
 - on CPU
 - minimum revision to the renderer
 - revision to the eval() function
- Storage
 - 1 million 2D texture samples with colors: 19 MB
 - 1 million 3D volume samples without color: 11 MB

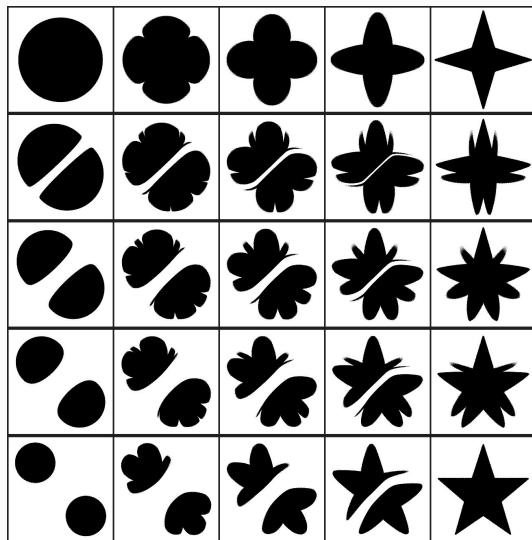


Results & Comparisons

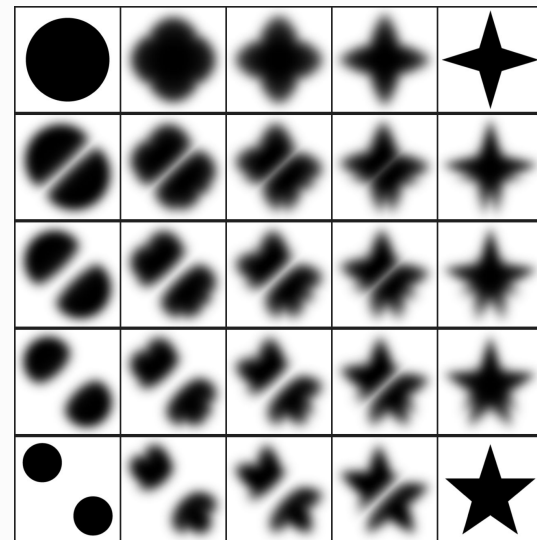


2D Texture Blending

- Comparison with Solomon et al.;
- Less runtime, and sharper results;
- Feydy et al. provides a sharper but different solver.



Ours. Pre.: 834.7s
Query: 7.3s

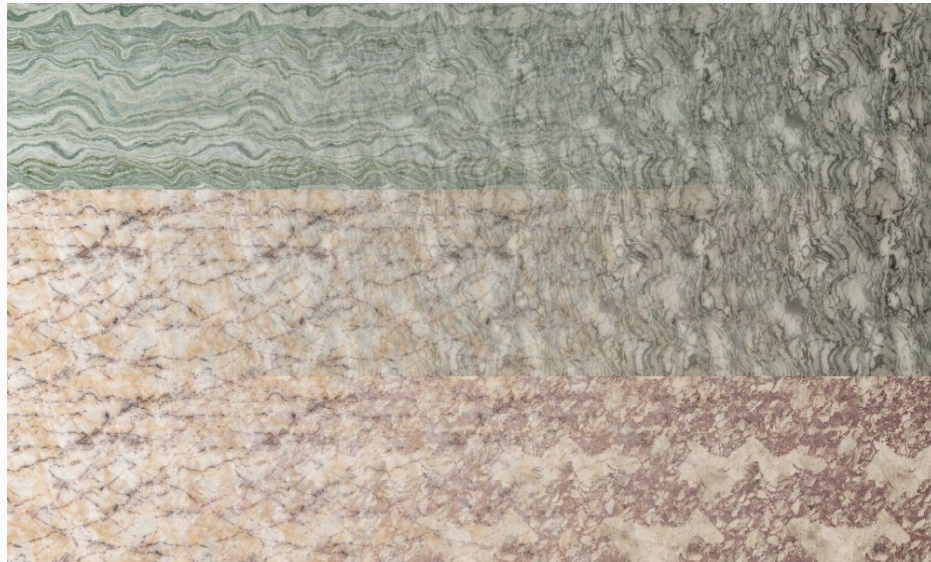


Convolutional Wasserstein
 $\epsilon = 0.005$, 21.2s

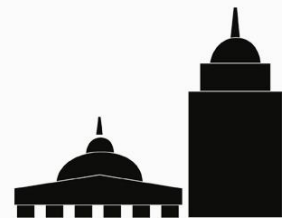


2D Tileable Texture Blending

- Inspired by Matusik et al.;
- Textures from Adobe Substance 3D Asset;
- Also included rendering results.

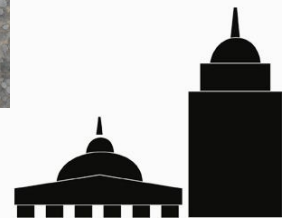
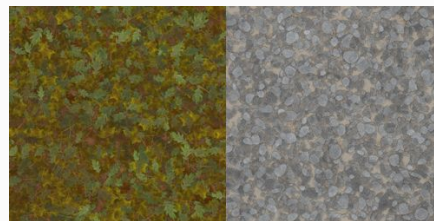
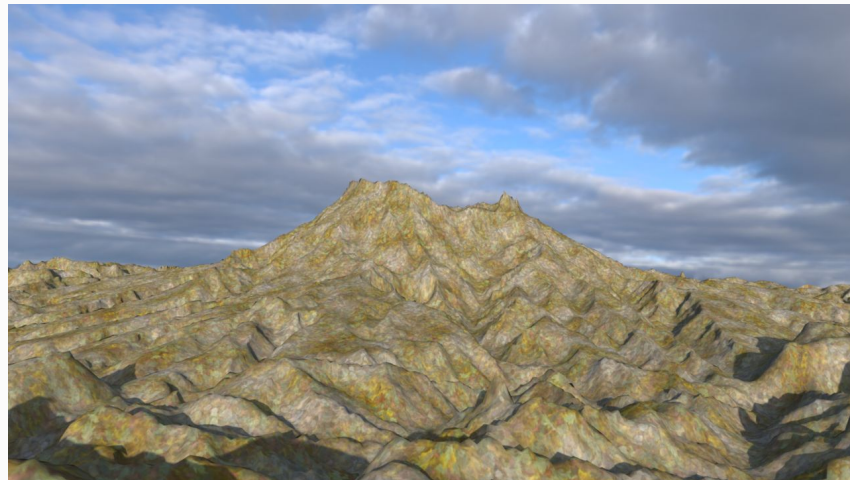


2D Tileable Texture Blending



Tileable Texture Blending with Perlin Noise

- Extension to tileable blending and query structure;
- Perlin noise provide the blending weight;
- Query structure allows focusing on blending a small range of data instead of generating the whole chunk of data;
- Seamlessly and infinitely extendable.



Tileable Texture Blending with Perlin Noise



Level-of-Detail (LOD) and Range Query

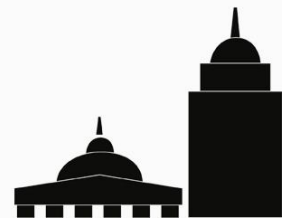
- LOD: detail decreases as the model moves away from the viewer;
- Range query: less traversal, less details, higher efficiency;
- Move the middle three textures away from the camera along the viewing direction.



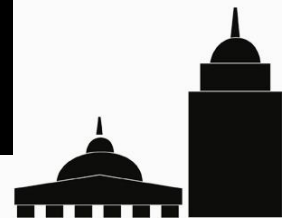
Closer view
512 spp, time = 33.17 mins



Further view
512 spp, time = 27.51 mins

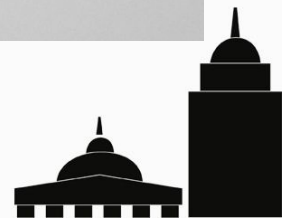


3D Volume 2-Way Interpolation



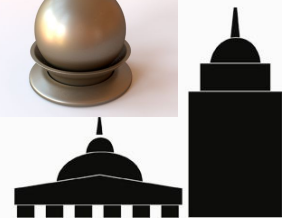
3D Volume 3-Way Interpolation

- 3-way volume blending inspired by Solomon et al.;
- Inputs are on the corners.



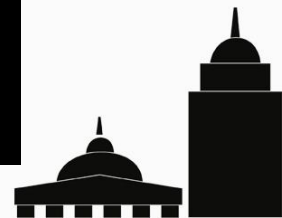
BRDF 4-Way Blending

- Measured BRDF from the MERL database;
- Inputs on the corners, and the blending weight starts at 1.0 for each BRDF on the corner and decreases by 0.2 with each step;
- Blending materials with different roughness, metallic, or Lambertian values.

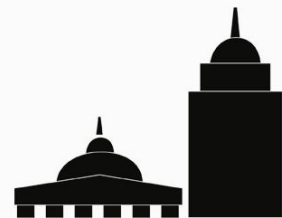


BRDF 4-Way Blending

Interpolating BRDF



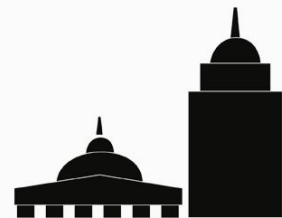
Conclusion & Limitations



Conclusion

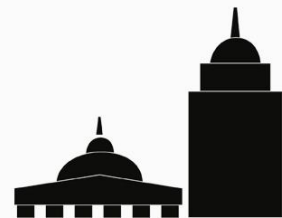
A **general**, efficient, and cross-dimensional **natural** appearance synthesis and rendering solution into traditional renderers.

- Precomputation:
 - OT calculate maps between a proxy distribution and each input distributions
- Runtime:
 - hierarchical point or range query structure
 - cross-dimensional multiple-way blending
 - generation of necessary data only



Limitations

- Lack a theoretical analysis about proxy selection;
- Lack a analysis about sample density analysis;
- Require further study of using proxy vs without proxy.



Thank you so much!

