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  title = {Joint Simulation of Transmission X-ray Imaging on {GPU} and Patient's Respiration on {CPU}},
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  annotation = {AAPM Annual Meeting, Jul~18--22, 2010},
  abstract = {Purpose: We previously proposed to compute the X-ray attenuation
from polygons directly on the GPU, using OpenGL, to significantly
increase performance without loss of accuracy. The method has been
deployed into a training simulator for percutaneous transhepatic
cholangiography. The simulations were however restricted to
monochromatic X-rays using a point source. They now take into account
both the geometrical blur and polychromatic X-rays.
Method and Materials: To implement the Beer-Lambert law with a
polychromatic beam, additional loops have been included in the
simulation pipeline. It is split into rendering passes and uses frame
buffer objects to store intermediate results. The source shape is
modeled using a variable number of point sources and the incident beam
is split into discrete energy channels. The respiration model is composed
of ribs, spine, lungs, liver, diaphragm and the external skin. The organ
motion simulation is based on anatomical and physiological studies:
the model is monitored by two independent active components:
the ribs with a kinematics law and the diaphragm tendon with an up and
down translation. Other soft-tissue components are passively deformed
using a 3D extension of the ChainMail algorithm. The respiration rate
is also tunable to modify the respiratory profile.
 Results: We have extended the simulation pipeline to take into account
focal spots that cause geometric unsharpness and polychromatic X-rays,
and dynamic polygon meshes of a breathing patient can be used as input data.
Conclusions: X-ray transmission images can be fully simulated on the GPU,
by using the Beer-Lambert law with polychromatism and taking into account
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# Joint Simulation of Transmission X-ray Imaging on GPU and Patient's Respiration on CPU

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## Purpose

We previously proposed to compute the X-ray attenuation from polygons directly on the GPU, using OpenGL, to significantly increase performance without loss of accuracy. The method has been deployed into a training simulator for percutaneous transhepatic cholangiography. The simulations were however restricted to monochromatic X-rays using a point source. They now take into account both the geometrical blur and polychromatic X-rays.

## Method and Materials

To implement the Beer-Lambert law with a polychromatic beam, additional loops have been included in the simulation pipeline. It is split into rendering passes and uses frame buffer objects to store intermediate results. The source shape is modeled using a variable number of point sources and the incident beam is split into discrete energy channels. The respiration model is composed of ribs, spine, lungs, liver, diaphragm and the external skin. The organ motion simulation is based on anatomical and physiological studies: the model is monitored by two independent active components: the ribs with a kinematics law and the diaphragm tendon with an up and down translation. Other soft-tissue components are passively deformed using a 3D extension of the ChainMail algorithm. The respiration rate is also tunable to modify the respiratory profile.

## Results

We have extended the simulation pipeline to take into account focal spots that cause geometric unsharpness and polychromatic X-rays, and dynamic polygon meshes of a breathing patient can be used as input data.

## Conclusions

X-ray transmission images can be fully simulated on the GPU, by using the Beer-Lambert law with polychromatism and taking into account the shape of the source. The respiration of the patient can be modeled to produce dynamic meshes. This is a useful development to improve the level of realism in simulations, when it is needed to retain both speed and accuracy.

# Joint Simulation of Transmission X-ray Imaging on GPU and Patient's Respiration on CPU

We present a simulation framework that combines respiratory motion and X-ray imaging. Our aim is to provide a validated training tool for interventional radiology to perform percutaneous transhepatic cholangiography (PTC). A CPU-based set of algorithms is presented to model the organ behaviour during breathing. Soft tissue deformation is computed with an extension of the ChainMail method and rigid elements move according to kinematic laws. A GPU-based surface rendering method is proposed to compute the X-ray image using the Beer-Lambert law. We demonstrate the efficiency of this approach by combining both visual and haptic cues and showing that interactive frame rates can be achieved.

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### Introduction

Introduction PTC consists in injecting a contrast agent into a bile duct within the liver to visualize the bilary tract using fluoroscopy (real-time X-ray images). The training of PTC is still performed as an apprenticeship within patients, and we have developed a virtual reality simulator to provide an alternative '. We present here the implementation of the two main software components of our simulator: i) the respiration modeling (inc. soft tissue deformation), and ii) the simulation of X-ray imaging.

**Respiration Modelling** ation model consists of a combination of i) rib motion nematic law, and ii) a translation of the diaphragm



Diaphragm heterogeneous concession. motion: motion: Tendon. They are respectively defined here by the angles u and  $\beta$ Wilson if t et al <sup>2</sup> measured these angles for five subjects : Functional Residual Capacity (FRC) and Total Lung Capaci (TLC). The rib rotation was thus decomposed into two distin rotations. We consider here that the rest position is at FRC. In on model, the rotation angles are given by the following equations:  $a' = (\sigma_{\rm TLC} - \sigma_{\rm FRC}) \times \frac{1}{2} \times (1 + \sin(2\pi ft))$   $\beta' = ((\beta_{\rm TLC} - \beta_{\rm FRC}) \times \frac{1}{2} \times (1 + \sin(2\pi ft))$ subjects a ing Capacity two distinct In ou

where: a' is the reduced 'pump handle' a(RC) = 0 and  $a'(TLC) = a_{TLC} - a_{REC} \beta'$  is the reduced 'bucket handle' angle  $\beta'(FRC) = 0$  and  $\beta'(TLC) = \beta_{TLC} - \beta_{REC} f$  is the respiration frequency and *i* is the time.

d r is the time. e central tendon is forced to have a sinusoidal movement alor e vertical axis similarly to the previous Equations. Duri natation, the central tendon has a downward movement, which morhorous with the expanding rib cage. As the links fort intral tendon are rigid, the points corresponding to this tendon v follow the exact same movement Other points of the diaphragm v corrected chas



ChainMail rules. However, the diaphragm is also supposed to follow the movement of the ribs. To do this, the idea is to "fix" some points of the diaphragm to some points of the diaphragm to some points of the diaphragm will be a combination of the downward movement of the diaphragm will be a combination of the downward movement of the central lendon and the movement induced by the contact points with the ribs.

#### Soft Tissue Deformation

and bones are modeled using rigid bodies. Other cal structures are modeled as soft tissues. Their tion is performed using the 3D ChainMail algorithm n proposed by Li and Brodlie<sup>3</sup>. Mesh elements are Ribs deformation extension pro

interconnected as links in a chain. So, within a certain limit, each point can move freely without influencing its neighbours. When one element of the object is moved and reaches this limit, the neighbours are forced to move in a chain reaction that is governed by the stiffness of the links in the mesh. The ChainMail parameters to simulate the deformation behaviour include the compression, the stretching and the shearing. The ChainMail parameters are set for each kind of tissue independently.

X-ray Imaging Simulation This dynamic data is used as input for our X-ray imaging simulation using the Beer-Lambert law. It relates the absorption of light to the properties of the material through which the light is traveling. For a monochromatic includer X-ray beam, it is:

### $N_{out}(E) = N_{in}(E) \exp \left(-\sum_{i=0}^{i < objs} \mu(E, i)L_p(i)\right)$

with  $N_{sl}(E)$  the number of incident photons at energy  $E, N_{sal}(E)$  the number of transmitted photons at energy  $E, \delta p_{ij}$  is the total number of objects in the 3D scene,  $p_i$  is the linear attenuation coefficient (in cm<sup>3</sup>), which depends on; i)  $E_i$  the energy of incident photons, and ii) the material properties of the object.  $L_{s}(p)$  is the path length (in cm) of the ray in the  $t^{\rm p}$  object.

 $L_p(i)$  is computed for each polygon mesh using the method proposed by Freud *et al* <sup>4</sup>. Normal vectors at the surface of objects are outward. The path length of the ray in a given object can be written as follows:  $L_p = \sum_i - (\operatorname{sgn}(\operatorname{viewVec} \cdot \mathbf{N}_i) \times d_i)$ 

where *i* refers to the  $\beta^{i}$  intersection found in an arbitrary order, *d*, is the distance from the X-ray source to the intersection point of the ray with the triangle, syn(twv ex-N) stands for the sign of the dot product between view Yee and N. ns can be efficiently achieved on the enGL Shading Language (GLSL) <sup>5</sup>. We show GPU us



#### Geometric unsharpness

The shape of the source is modeled using a varial point sources <sup>6</sup>. Each point is assigned a fraction number of photons in the system. number of of



Pressure and the provided and the pro



Conclusion The respiration of the patient can be modeled to produce dynamic meshes in real-lime. This data is then used to simulate X-ray transmission imaging on the GPU, by using the Beer-Lambert law with polychromatism and taking into account the shape of the source. This is a useful development to improve the level of realism in simulations, when it is needed to retain both speed and accuracy. n simula accuracy.

FREUD N., DUVAUCHELLE P., LÉTANG J. M., BABOT D.: Fast and robust ray casting porthims for virtual X-ray imaging. Nucl Instrum Methods Phys Res B 248, 1 (2006), 175-ER M., FREUD N., LÉTANG J. M., JO J. In Proc Theory Pract Comput Graph NIER M., FREUD N., LÉTANG J. N I., JOHN N. W.: Simulation or Array raph (2009), pp. 25–32. J. M., JOHN N. W.: Accelerated Partware In EG 2010

