MASARYK UNIVERSITY FACULTY OF INFORMATICS



Maya2CellVIEW: 3D Package Integrated Tool for Creating Large and Complex Molecular Scenes

Master's Thesis

David Kouřil

Brno, Fall 2016

MASARYK UNIVERSITY FACULTY OF INFORMATICS



Maya2CellVIEW: 3D Package Integrated Tool for Creating Large and Complex Molecular Scenes

Master's Thesis

David Kouřil

Brno, Fall 2016

This is where a copy of the official signed thesis assignment and a copy of the Statement of an Author is located in the printed version of the document.

Declaration

Hereby I declare that this paper is my original authorial work, which I have worked out on my own. All sources, references, and literature used or excerpted during elaboration of this work are properly cited and listed in complete reference to the due source.

David Kouřil

Advisor: Ivan Viola

Acknowledgement

This is the acknowledgement for my thesis, which can span multiple paragraphs.

Abstract

Scientific illustrators communicate the cutting edge of research through their illustrations. There are numerous software tools that assist them with this job. The aim of this thesis is to push abilities of illustrators working on a large scale molecular scenes. This is done by connecting two software packages - Maya and cellVIEW - combining the rendering possibilities of cellVIEW and modeling tools of Maya which results in more effective and efficient workflow.

Keywords

keyword1, keyword2, ...

Contents

| 1 | Introduction | 1 |
|---|---|----|
| | 1.1 Biological visualization and illustration | |
| | 1.2 Motivation | 1 |
| 2 | State of the art | 3 |
| 3 | OLD - Overview of the two programs that I'm going to be | |
| | bridging - TODO better title | 5 |
| | 3.1 <i>CellVIEW</i> | 5 |
| | 3.1.1 Unity | 5 |
| | 3.2 <i>Maya</i> | 5 |
| | 3.2.1 Extensibility | 6 |
| 4 | Method | 7 |
| 5 | Implementation | 9 |
| | 5.1 <i>WinAPI</i> | 9 |
| | 5.2 <i>Maya side</i> | 9 |
| | 5.3 <i>Unity side</i> | 9 |
| 6 | Demonstration | 11 |
| 7 | Discussion, future work | 13 |
| A | An appendix | 15 |

List of Tables

List of Figures

1 Introduction

In this age, scientists come to new findings everyday. Unfortunately, not all of these are ever shown to the general public. It could be because of several reasons. One of them might be that the new facts are not easily visualizeable. This can also mean that the things that they found out are a piece of a bigger picture. This is the job of scientific illustrator. These artists are usually experts in their own scientific domain and what they do, is they try to visualize the science in their domain with easily understandable images, animation and other media.

2 Introduction - old

2.1 Biological visualization and illustration

Here I should somehow present the field (but I don't know what field exactly - biological visualization, molecular visualization?)

2.2 Motivation

In this thesis we focused on providing better tools for scientific illustrators. The idea was to bridge software that they are used to (represented by Maya) and software that brings modern rendering abilities to the domain of molecular rendering (represented by CellVIEW). Emphasize the difference between offline and realtime rendering.

3 State of the art

In this chapter we will look at the current tools for authoring of scientific illustrations. Illustrators have gone a long way from doing these works by hand and nowadays there exist a variety of programs that help them do their job communicating the scientific findings. Barely any of these tools is perfect. Some of the tools are very closed and tied to it's creator. The ideal case would be if any new user could approach any of these tools and with a little training be able to use it. This is not the case, unfortunately. From the ones that are fairly simple from the user experience stand point is Molecular Flipbook[ref]. It provides user the ability to create simple molecular models and animate them with a keyframe-like animation workflow.

An important part of any scientific illustration are the data the one is trying to illustrate. In the case of molecular models, there is cellPACK which helps us with this. cellPACK is a software that assembles large molecular scenes from a description (a recipe) of how this scene should look, what it should contain and in what quantity.

3.1 Maya

Autodesk Maya is not a tool for molecular visualization per-se.

3.2 cellVIEW

4 OLD - Overview of the two programs that I'm going to be bridging - TODO better title

Here I will talk about in which context are these two 'systems' used and what is each for. The point of this project was to make a bridge between two programs. In this chapter we will look at both of them, see what they bring to the table when used separately, what they lack and how can be benefit from connecting them in this way.

4.1 CellVIEW

Briefly describe what CellVIEW is about. Give links to the papers that are describing CellVIEW and those that are using it.

4.1.1 Unity

I don't know if this is needed. I could just go little bit into detail of why is Unity good (which is funny now that we are probably going away from it)

4.2 Maya

Point to examples of animations that are produced with Maya. Maya is a 3D content authoring software package developed by Autodesk. It has been introduced in 1998 and since than it has grown into defacto industry standard in many fields that use computer graphics (most notably movies and computer games industry). Because of it's versatility, Maya has also been used for creating scientific illustrations. Although Maya is a very robust piece of software that gives it's users ability to create various types of 3D scenes, it isn't always the best tool for the job. There exist other software packages that might be more specialized and better suited for a task at hand.

4.2.1 Extensibility

Luckily, Maya comes with an enormously useful ability to be extended with so called plugins. Autodesk provides an API that programmers can use to extend Maya's functionality. Several plugins have been developed specifically for the field of molecular visualization and illustration. As an example, we can name Molecular Maya which brings users the ability to import molecular data from online Protein Data Bank. The plugin then allows the user to choose how this data should be represented in a 3D scene and Maya then takes care of rendering such structure.

5 Method

I have introduced the field. I have sketched the problem that we are trying to solve (improving the workflow of animators/illustrators). Now it's time to describe the suggested solution. We talked about improving the workflow of scientific illustrator that uses Maya as an authoring software. The way that we are going to accomplish this is by taking advantage of extensibility of both Maya and Unity.

6 Implementation

6.1 WinAPI

6.2 Maya side

Topics - Architecture of the plugin, what parts/classes of API are used, memory layout, what I tried and didn't work.

6.3 Unity side

Topics - Architecture of the plugin, very generally about plugins (it's just a basic C++ dll plugin), interface between C++ and C#

7 Demonstration

In this chapter we present two use cases of how the user can approach illustration with this new proposed system.

8 Discussion, future work

What optimizations I could implement, how it could be used and what would be needed to implemented for that.

A An appendix

Here you can insert the appendices of your thesis.