



EDINBURGH NAPIER UNIVERSITY

SET09119 PHYSICS-BASED ANIMATION

Workbook

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September 16, 2016

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Version 1.3, 3 November 2008

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Chapter 1

Getting Started

1.1 Welcome

Welcome to Physics-Based Animation! This module and accompanying workbook will guide you through the process of simulating the vast mechanisms of real-world physics within software. Expect collisions, explosions, particles, and plenty of cubes and spheres rolling around everywhere.

The pre-requisite knowledge you require is:

- A Strong grasp of mathematical concepts such as trigonometry, algebra, and geometry. Linear algebra and matrix mathematics are crucial skills for this module.
- An adept or above level of skill with the C++ language.
- Knowledge of Computer Graphics technology and theory.
- Strong software debugging skills
- A willingness to spend time solving the problems presented in this workbook. Some of the exercises are challenging and require effort. There is no avoiding this. However, solving these problems will significantly aid your understanding.

1.2 Workbook Style

The workbook is task and lesson based, requiring you to solve problems to complete the code provided as a starting point for the lesson. This code will either be in the main application source file (`main.cpp`) for the lesson, or in one or more shader files (typically found in the `resources/shaders` folder). The sections you have to complete are highlighted in stars:

```
1  \\ *****
2  \\ You have to complete this problem
3  \\ *****
```

The problems faced generally build on previous lessons until you become familiar with the work involved.

1.3 Physics Framework

This module provides example code for you to build upon throughout the different topics. To make life easier, a small physics library has been included, this contains some useful helper functions and abstracts most of the graphics logic away from the main tutorial code. This physics library is built on top of the Napier graphics framework, which you should be familiar with. The purpose of this module is to teach physics, not graphics, however; the two are closely linked so you may need to dive down into the libraries to write/modify some graphics code.

1.3.1 Libraries

Other than the libraries used by the graphics framework (GLEW, GLFW, ASSIMP...) the physics code makes use of the following libraries:

GLM the GL Mathematics library. GLM is also cross-platform and a header only library (it has a number of different headers we include for different purposes). GLM provides the mathematical types and functions required to work with OpenGL (which provides no basic mathematical types).

IMGUI , a small graphical user interface library for C++, is particularly suited to integration in realtime 3D applications. It favours simplicity and productivity and is easy to implement and use.

1.4 Getting the Code - Clone The Repository from GitHub

Everything you need to get started (including this workbook) is hosted in a git Repository. You will need to clone your own copy to work with.

```
1 git clone https://github.com/NapierUniversity/enu_pba.git
```


1.5 Configuring the code - CMake

1.5.1 Intro to CMake

The physics project makes use of the CMake build system, this allows for cross-platform compatibility, dependency acquisition and general purpose code housekeeping. If you are new to CMake, here is a simple explanation of its purpose.

CMake uses a configuration script to generate Make Files (or IDE Project files) for your project. Instead of shipping a makefile which would have to be updated, and would be specific to one platform, only the project code and a CmakeLists.txt config script is needed. The same goes for Visual studio Solution files, which are large, difficult to version, and can only be used by Visual studio.

Instead of maintaining a visual studio sln file, CMake will generate one for you. CMake does not actually *build* your code, it just *configures* your build environment for you so you can build the software.

CMake is also used to pull down the additional libraries we will need for this project. This means that you are getting a fresh and up-to date copy of all the libraries, and you will be compiling them specifically for your system. Say goodbye to linker and VCRuntime errors.

1.5.2 Running CMake

To keep the source directory (the directory pulled from Git) clean, we tell CMake to generate our build files in a separate directory, This is called an Out of Source Build. We can do this all from the command line easily:

```
1 mkdir physics_build
2 cd physics_build
3 cmake -G "Visual Studio 14 2015 Win64" ../enu_pba/
```

CMake Has a GUI interface also, this is useful if you are configuring new software and need to see and set options. This project has all the settings set to sensible defaults so you should not need to touch anything other than the configure and generate buttons. The steps are Shown in Figure 1.1.

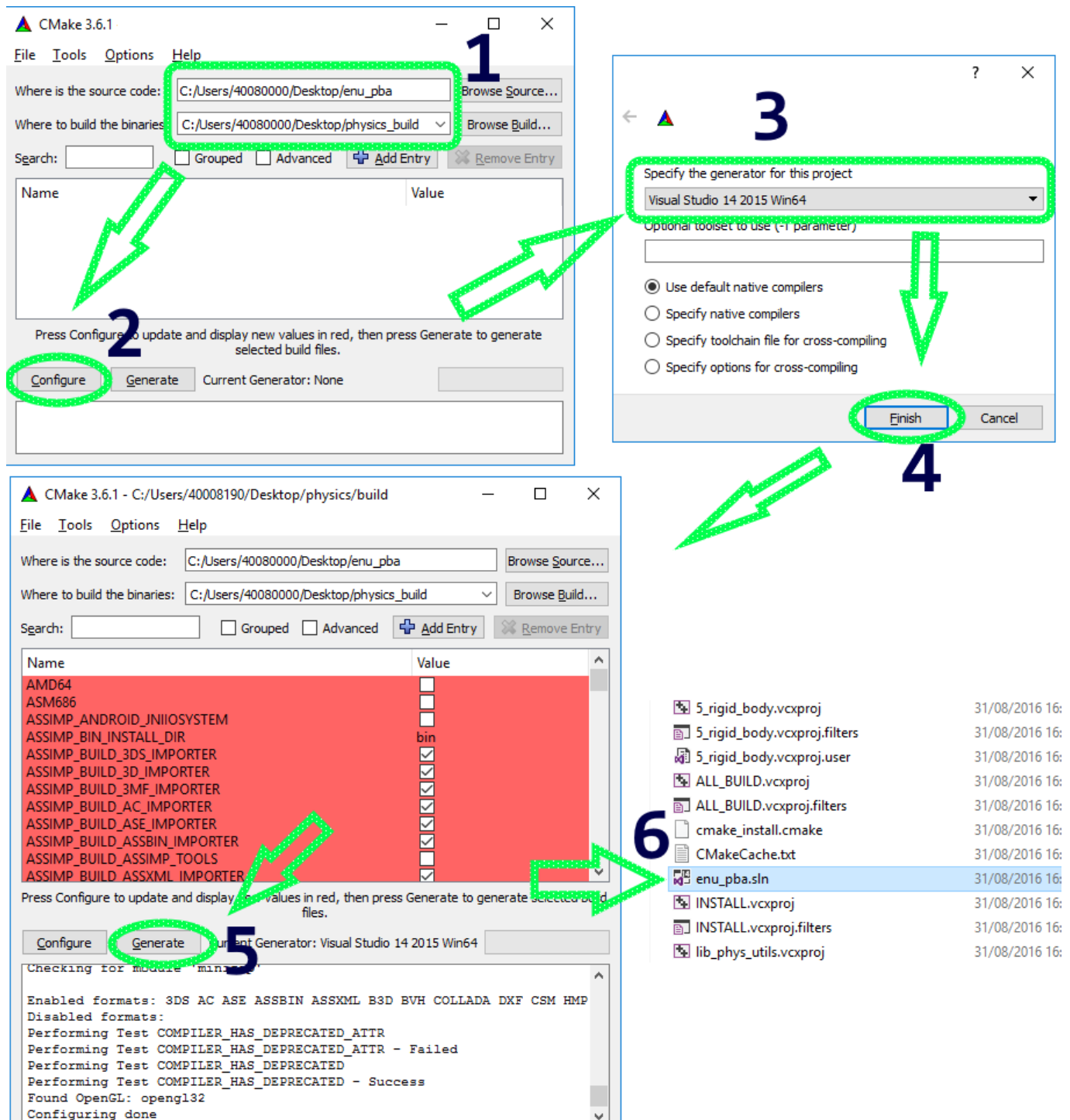


Figure 1.1: CMake GUI Build Steps

1.6 Building the code - Visual Studio

Once CMake has created the Solution file and you have opened it, you should see the following projects:

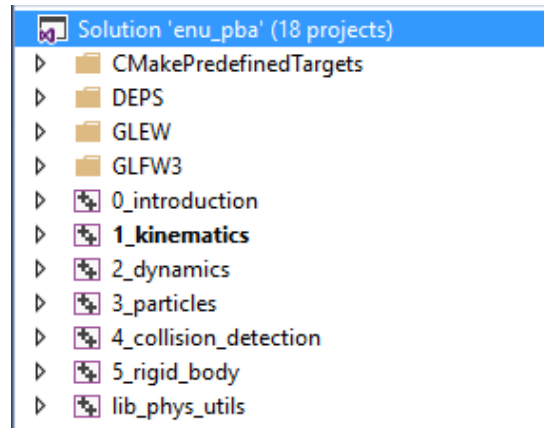


Figure 1.2: Visual Studio projects

The projects within the sub-folders are the libraries which we will use, so you don't need to pay attention to them. The CMakePredefinedTargets contains some CMake functions, specifically useful is the ZERO_CHECK project, this will trigger a CMake configure and rebuild without having to leave VS. This is useful if anything in the source directory has changed.

1.7 Running the Code

You should run this application to see the output. You will get a window as shown in Figure 1.3.

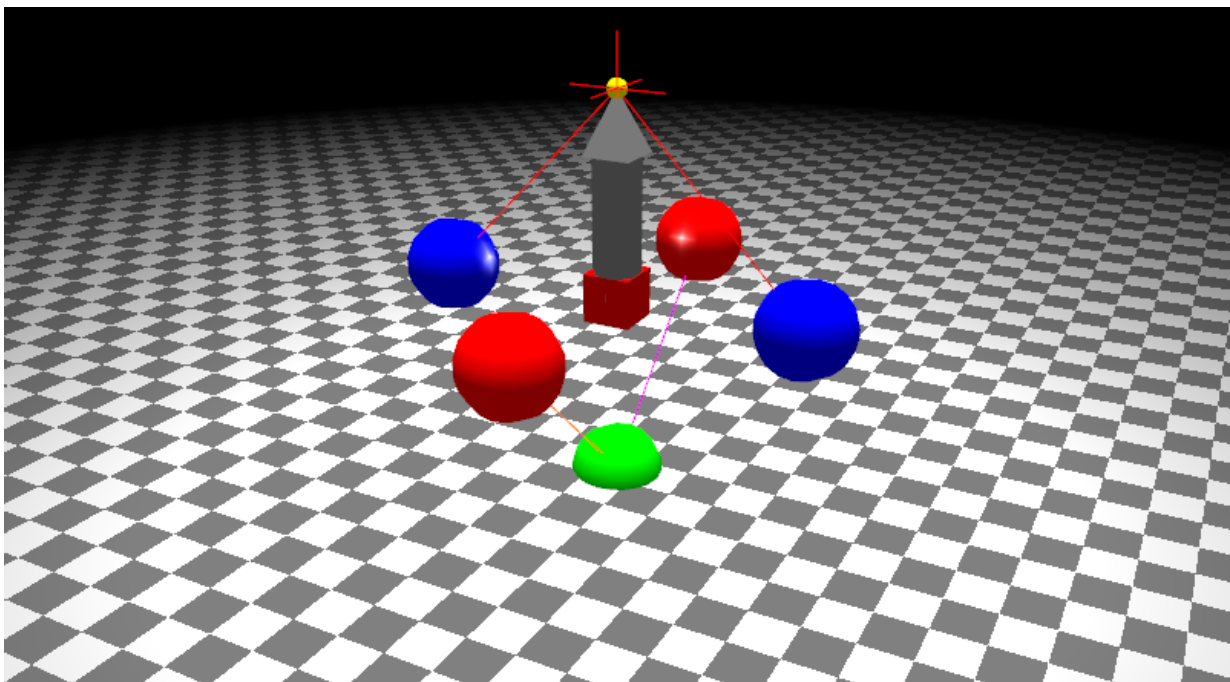


Figure 1.3: Output from Basic Application

Chapter 2

Inverse Kinematics

2.1 Introduction

The topic of this practical is the implementation of a real-time inverse kinematic simulation (i.e., a linked chain of rigid body objects). The user should be able to interact with the simulation while it is running (e.g., through the mouse or keyboard) to control the inverse kinematic target end-effector. The inverse kinematic chain can be implemented using either the cyclic coordinate descent (CCD) algorithm or the Jacobian pseudo-inverse matrix method.

Tasks

1. Visually display interconnected chain of 3D rigid bodies (i.e., base and end-effector)
2. Implement an uncomplicated single-joint IK system using an analytical method (i.e., axis-angle) which follows the mouse cursor around the screen (Section 2)
3. Implement a linked-chain of interconnected limbs using an iterative inverse kinematic technique (e.g., CCD IK with details given in Section 3)
4. User input (e.g., mouse or keyboard) to control and move the end-effect target
5. Areas to explore, include, altering the number of links (e.g., 10 to 1000). Adding physical constraints (i.e., limit the joint angles +/- specified amount). Modify the code to support multiple end-effectors (e.g., tree-like structure)

2.2 Overview

Principles and Concepts The application of kinematic algorithms for animation is used to control articulated postures based on a simple definition of joint angles and limb lengths. These structures may take practically any form from a humanoid bipedal character to just about anything that can be imagined using a hierarchy.

Hierarchy Structures described using a hierarchical form are defined using a parent-child system similar to that of tree structure. In the case of computer characters, each rigid limb of the structure is a child node in the tree whose parent node provides a reference point from which it is described. The parents are themselves child nodes of limbs above in the hierarchy and this recursive relationship

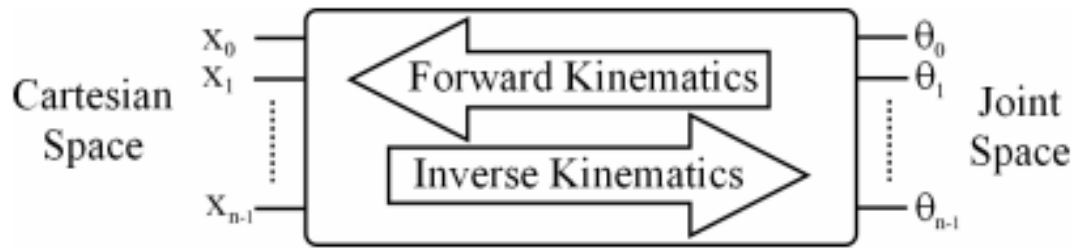


Figure 2.1: **Forward & Inverse Kinematics** Illustrating the relationship between forward and inverse kinematics parameters

continues up to a root node (i.e., the bottom of the tree). The parent of the root node is effectively taken as the global frame of reference and defined as such.

End-Effectors At the top end of the tree are leaf nodes which are children that have no descendants of their own. An end-effector in terms of kinematic chains is any node within the hierarchy that an animator wishes to directly position, for example, to interact with the environment. End-effectors are commonly the leaf nodes of an articulated structure, such as, the feet and hands of an animated character since they generally interact with the world.

Graphics Inverse kinematics is a programmatic solution for controlling the animation of rigid models. When artists generate an animated model for a virtual environment, such as a game, the animations created by the artists won't necessarily work for every position in the world. For example, if an artist were to create an animation of a character pressing a button, the animation would only work so long as the button was always at the same position relative to the model. If the button were to be moved up or down the pre-made animation would have no way to account for that. With inverse kinematics, we can reconfigure the interconnected set of links (e.g., animate the model) so the end-effector (e.g., a hand) can be placed anywhere.

Inverse Kinematic (IK) Methods

1. Analytical (e.g., Geometric Analysis)
2. Heuristic (e.g., CCD)
3. Algebraic Solver (e.g., Jacobian & Gradient-Based Search)

2.3 Analytical Inverse Kinematics

The geometric/analytical algorithms tend to be very quick because they reduce the IK problem to a mathematical equation that need only be evaluated in a single step to produce a result (e.g., see Figure 2.2). The limitations of this class of solver becomes apparent in the case of large chains. In such cases, the task of reducing the problem to a single-step mathematical equation is impractical. Therefore geometric/analytical techniques tend to be less useful in the field of character animation.

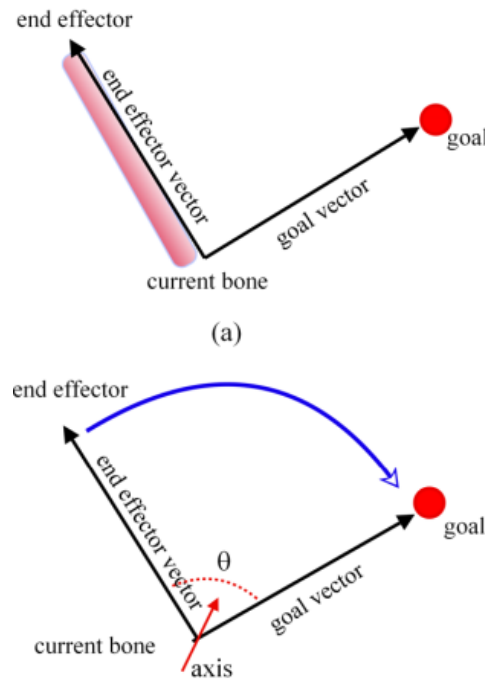


Figure 2.2: **Analytical Inverse Kinematic Example**Single limb and joint angle (solved using the dot & cross product)

2.4 Cyclic Coordinate Descent (CCD)

IK solvers that are based on CCD use an iterative approach that takes multiple steps towards a solution. CCD works by analysing each joint one-by-one in a progressive refinement philosophy. Starts with the last joint in the chain (e.g., a hand for a character) and tries to rotate it to orientate it toward the target. The steps that the method takes are formed heuristically, therefore each step can be performed relatively quickly. An example of a possible heuristic would be to minimise the angle between pairs of vectors created when projecting lines through the current node and end-effector and current node and desired location. However, because the iterative step is heuristically driven, accuracy is normally the price paid for speed. Another issue with this technique is that one joint angle is updated at a time as opposed to the complete hierarchical structure. This has the undesirable and unrealistic result of earlier joints moving much more than later limbs in the IK chain.

Implementation The Cyclic Coordinate Descent (CCD) algorithm is an iterative IK solution. The basic idea of the CCD algorithm is to loop over each bone in the IK chain and rotate it such that the end effector (which is typically the last bone in the chain) will move as close as it can to the final position

```

while (distance from end effector to target  $\geq$  threshold)  $\wedge$  (numloops < max) do
    Take current bone;
    Build vector V1 from bone pivot to end effector;
    Build vector V2 from bone pivot to target;
    Get the angle between V1 and V2;
    Get the rotation direction;
    Apply a differential rotation to the current bone;
    if current bone is the last bone in the chain then
        | restart at first bone;
    else
        | current bone is the next bone in chain;
    end
end

```

Algorithm 1: Algorithm for the CCD system

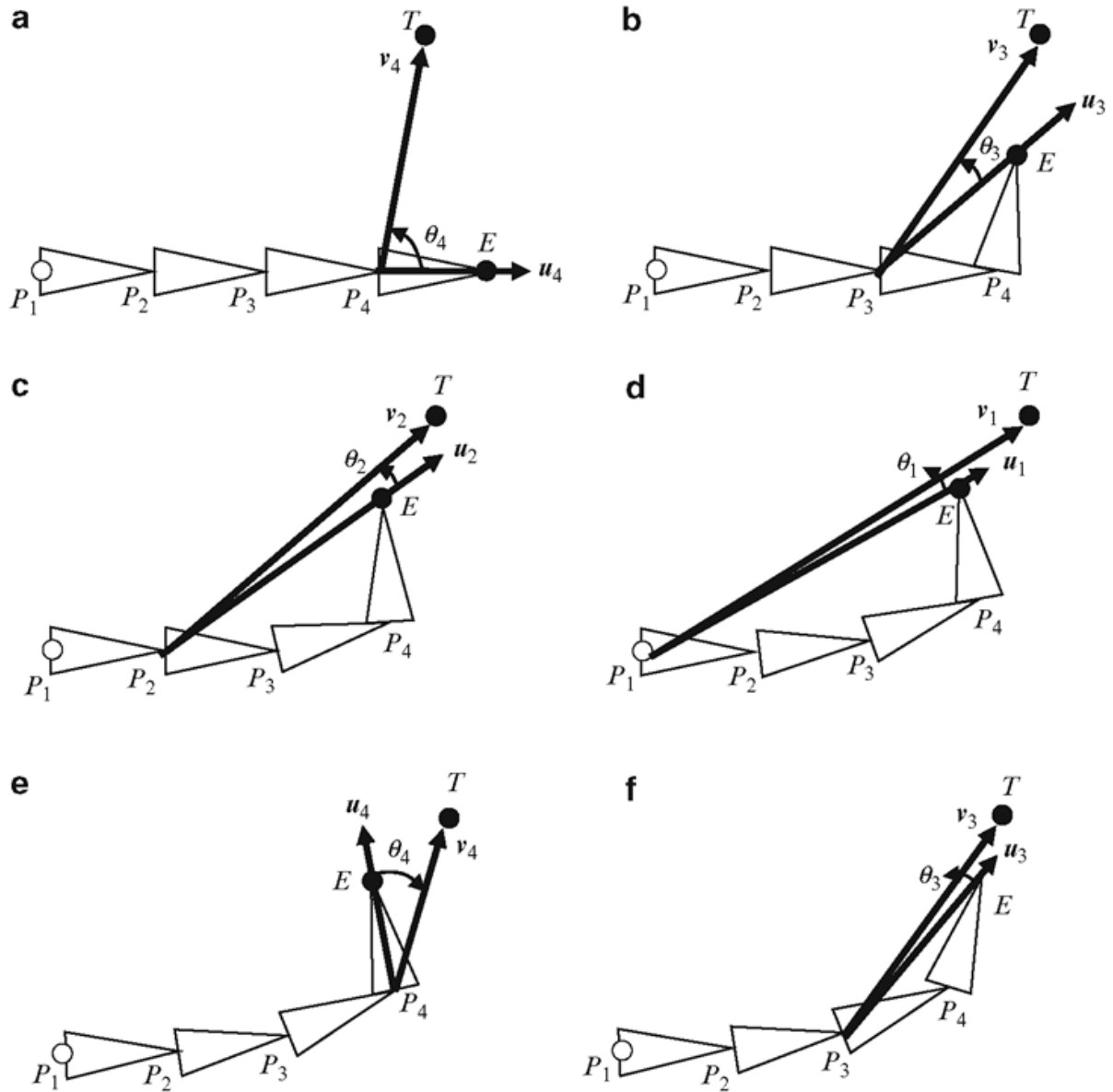


Figure 2.3: Sequence of rotations performed by CCD algorithm on a 4-link joint chain[1]

Figure 2.3 shows the principle of the CCD technique. (a) Shows The initial state

(b) The end effector joint P_4 has been moved to bring the vector U_4 as close as possible to V_4 , the next vector for the next joint P_3 is then calculated.

(c) The joint P_3 has been moved to bring the vector U_3 as close as possible to V_3 , this brings the end effector E close to the target T

(d) The Process repeats for the last joint

(e..f) The process restarts from the end joint P_4

2.5 Iterative Jacobian Pseudo Inverse

Due to their scalability, numerical techniques often form part of an inverse kinematics solver. However, because of their iterative nature, such methods can be slow. So far research into the field of kinematics has failed to find a general nonnumerical solution to the problem. Many researchers have proposed hybrid techniques yet these still rely on a numerical aspect. It is therefore important to find ways of using numerical techniques as efficiently as possible. In this paper we take a look at the Jacobian-based IK solver and techniques that allow this method to be used as an efficient real-time IK solver.

The end effector velocities are related to the joint velocities of the robot through the Jacobian matrix as follows:

$$\dot{e} = J\dot{\theta}$$

Where J is the Jacobian representing the partial derivatives for the change in end-effectors locations with change in joint angles. Typically, the system of joints consists of a non-square Jacobian, hence, we need to perform the pseudo inverse operation to yield the joint velocities and solve for the angular displacement:

$$J^+ = J^T(JJ^T)^{-1}$$

$$\dot{\theta} = J^+\dot{e}$$

Where we are able to connect $\dot{\theta}$ for change in joint angles with change in end-effector location \dot{e} . This method sets the angle values to the pseudo inverse of the Jacobian. It tries to find a matrix which effectively inverts a non-square matrix. It has singularity issues which tend to the fact that certain directions are not reachable. The problem is that the method first loops through all angles and then needs to compute and store the Jacobian, perform a pseudo inversion, calculate the changes in the angle, and last apply the changes.

```

while distance from end effector to target  $\geq$  threshold do
    Compute  $J(e, \theta)$  for the current pose;
    Compute  $J^{-1}$  i.e., invert the Jacobian matrix;
     $\Delta e = \beta(t - e)$  - pick approximate step to take;
     $\Delta \theta = J^{-1}\Delta e$  - compute change in joint DOFs;
     $\theta_{current} = \theta_{previous} + \Delta \theta$  - apply change to DOFs;
end

```

Algorithm 2: Jacobian System

2.5.1 Calculating the Jacobian Matrix

The Jacobian J is a matrix that represents the change in joint angles $\Delta\theta$ to the displacement of end-effectors Δe . Each frame we calculate the Jacobian matrix from the current angles and end-effectors. We assume a right-handed coordinate system. To illustrate how we calculate the Jacobian for an articulated system, we consider the simple example shown in Figure 2.4

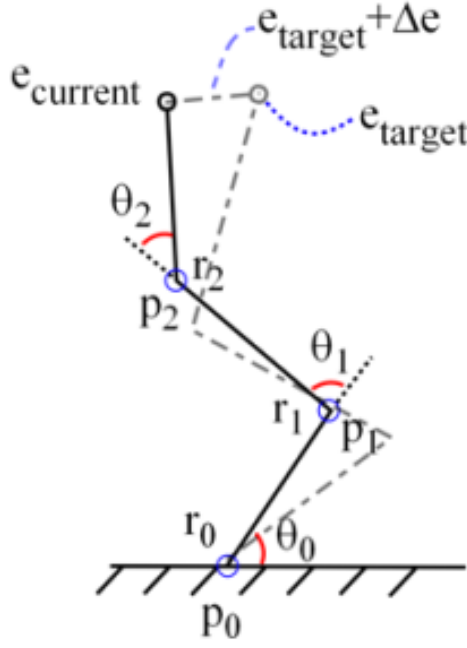


Figure 2.4: **2D 1DOF Jacobian IK Example** - Relationship between multiple joint angles and end-effectors.

Figure 2.4 demonstrates how we decompose the problem and represent it as a matrix for a sole linked chain with a single three degree of freedom (DOF) end-effector. We then extend this method to multiple linked-chains with multiple end-effectors (each with six DOF) to represent a structured hierarchy

$$\theta = \begin{bmatrix} \theta_0 \\ \theta_1 \\ \dots \\ \theta_n \end{bmatrix} \quad e = \begin{bmatrix} e_{current_x} \\ e_{current_y} \\ e_{current_z} \end{bmatrix}$$

where θ is the rotation of joint i relative to joint $i - 1$, and e for the end-effectors global position. The angles for each joint and the error for each end-effector are represented by matrices. From these matrices, we can determine that the end-effectors, and the joint angles are related. This leads to the forward kinematics definition, defined as:

$$e = f(\theta)$$

We can differentiate the kinematic equation for the relationship between end-effectors and angles. This relationship between change in angles and change in end-effectors location is represented by the Jacobian matrix.

$$\dot{e} = J(\dot{\theta})$$

The Jacobian J is the partial derivatives for the change in end-effectors locations by change in joint angles.

$$J = \frac{\delta e}{\delta \theta}$$

If we can re-arrange the kinematic problem:

$$\theta = f^{-1}(e)$$

We can conclude a similar relationship for the Jacobian:

$$\dot{\theta} = f^{-1}(\dot{e})$$

For small changes, we can approximate the differentials by their equivalent deltas:

$$\Delta e = e_{target} - e_{current}$$

For these small changes, we can then use the Jacobian to represent an approximate relationship between the changes of the end-effectors with the changes of the joint angles.

$$\Delta\theta = J^{-1}\Delta e$$

We can substitute the result back in:

$$\theta_{current} = \theta_{previous} + \Delta\theta$$

The practical method of calculating J in code is used:

$$\frac{\delta e}{\delta \theta_i} = r_j \times (e_{target} - p_j)$$

Where r_j is the axis of rotation for link j , e_{target} is the end-effectors target position, p_j is end position of link j

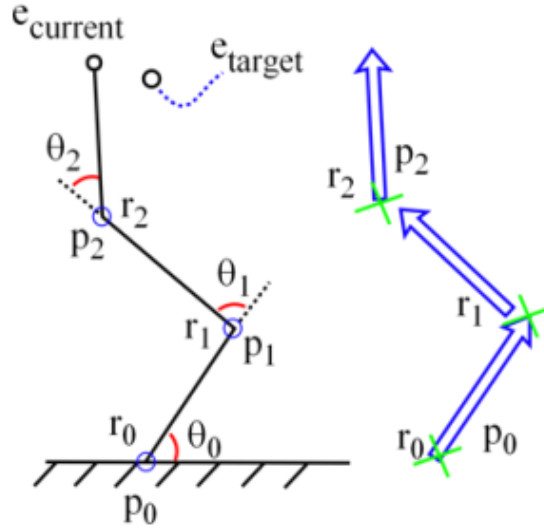


Figure 2.5: **Jacobian**- Iteratively calculating the Jacobian on a frame by frame basis.

For example, calculating the Jacobian for Figure 2.5 gives:

$$J = \begin{bmatrix} \frac{\delta e}{\delta \theta_0} \\ \frac{\delta e}{\delta \theta_1} \\ \frac{\delta e}{\delta \theta_2} \end{bmatrix} = \begin{bmatrix} r_0 \times (e_{current} - p_0) \\ r_1 \times (e_{current} - p_1) \\ r_2 \times (e_{current} - p_2) \end{bmatrix}$$

$$e = e_{current} - e_{target}$$

The Jacobian matrix is calculated for the system so that we can calculate the inverse and hence the solution. Once the Jacobian system has been defined, we iteratively solve for the change in angle using an approximate technique, such as, Pseudo-Inverse Transpose.

2.6 Summary

We have introduced the basic inverse kinematic concepts, and had a quick refresher course of the kinematics required to develop the algorithms. Your module work should now be structured in such a way that the inverse kinematic simulation is separate from the renderer update, and so that the implementation which you will develop can easily interface with the rest of your system (i.e., modular component programming).

2.7 Tutorial Exercises

Basic Exercises

1. Complete the 3 degrees of freedom example (ik_3dof.cpp)
2. Complete the 1 degree of freedom example (ik_1dof.cpp)
3. Add the option to only move the arm once a solution has been found, Lerp for smoothness. Careful! make sure you add a timeout/max attempts clause
4. Experiment with algorithmic enhancements to see if you can improve the process

Advanced Exercises

- Implement the FABRIK algorithm, as described in the paper [2] and [3]
- Implement an IK solver using the inverse Jacobian method

Chapter 3

Dynamics

3.1 Time

One of the main difference between working out real physics on paper and simulating an interactive simulation is that the world can change and thus the end state is not known in the simulation. For example: given the simple problem of how far will a projectile fly, or how quickly will a box fall, assuming you have all the required parameters (mass, gravity, distances and dimensions), then you can find the definite solution and be 100% accurate. When simulating a rocket flying, or a box falling in a physics engine, there is nothing stopping us using the same equations, finding out the same answer and then interpolating over time. Alas, it's not that simple, for at least two major reasons:

1. The player can alter the parameters during the flight of the box or rocket; they could put something in the flight path, or shoot the box.
2. If we only calculate the end state, simply interpolating between the start and end state will not show certain details. Items in the real world tend to accelerate and decelerate over time, interpolating would not show this accurately.

Fundamentally to build a simulation of real world physics, we must do exactly that: Build a simulator, not a calculator. However, the environment in which our simulator will run (on a CPU) has many small but significant differences to real world physics. The primary issue we will face is time. Relativity aside, time in the real world is continuous, but the time we have in our simulation happens in intervals.

Using some baseline numbers for an average game engine, we usually have 16 milliseconds to do everything we need to calculate the gamestate at one given point in time. After we have done these calculations, we move onto the next 'frame', and then calculate everything for a world that is now 16 milliseconds further on.

This can be a complex concept to grasp, but a crucial one. Every physics 'Tick' we must calculate where everything in the world must be at a single point in time. What this point in time actually is will depend on a few factors. To make matter more complex, games will never run at a constant frame-rate, so how many time the physics must 'tick' to keep pace with an accurate time-line must be well thought through. This section will cover the engineering challenging of deciding when and how much to "Tick"

3.1.1 Simple FPS-Fixed Timestep

The simplest method for when to run a physics tick would be once, every frame. We keep an accurate measurement on the "real time" (variable `t`), by using the delta-time passed into us.

```

1 double t;
2 void render() {...}
3
4 void physics_tick(time_now, delta_time){...}
5
6 void update(delta_time) {
7     do_game_logic();
8     physics_tick(t, delta_time);
9     t += delta_time;
10 }
```

This would be fine for only the most rudimentary of physics applications, one tick per frame is not nearly enough, so we can extend this method by doing multiple ticks per frame.

```

1 #define ticks_per_frame 10
2 void update(delta_time) {
3     do_game_logic();
4     for(i < ticks_per_frame)
5         physics_tick(t, delta_time / ticks_per_frame);
6     t += (delta_time / ticks_per_frame);
7 }
8 }
```

This would work, but we are at the mercy of the framerate. Imagine an extreme example where the renderer took ten minutes to kick out a frame, with our code above, we would be doing one physics tick per minute. Clearly this would cause issues, and so we must use a method that places a limit on the maximum delta-time we use for physics.

3.1.2 Maximum Timestep

```

1 #define max_physics_tick (1.0 / 600.0)
2 double t;
3 void render() {...}
4
5 void physics_tick(time_now, delta_time){...}
6
7 void update(delta_time) {
8     do_game_logic();
9     double remainingTime = delta_time;
10    while (remainingTime > 0.0) {
11        //never have a dt > max_physics_tick
12        double dt = glm::min(remainingTime, max_physics_tick);
13        remainingTime -= dt;
14        t += dt;
15        physics_tick(t, dt);
16    }
17 }
```

Here we define a maximum limit to DT (in this case: 600 fps), other than a physics stall (covered later) this method is much more stable and less prone to strange fluctuations in framerate. Having a DT that can vary *might* cause some issues with a physics simulation, and it *may* also be difficult to tune for performance. Generally there will be a sweet spot with the number of Ticks or magnitude of DT; having both vary might make life difficult, an issue the next method solves.

3.1.3 Fixed Timestep, non-fixed tick count

```
1 #define physics_tick_rate (1.0 / 60.0) / 10
2 double t;
3 void render() {...}
4
5 void physics_tick(time_now,delta_time){...}
6
7 void update(delta_time) {
8     do_game_logic();
9     static double accumulator += delta_time;
10    while (accumulator > physics_tick_rate) {
11        accumulator -= physics_tick_rate;
12        t += physics_tick_rate;
13        physics_tick(t, physics_tick_rate);
14    }
15 }
```

This is a subtle change from the previous Maximum Timestep method, this method uses a constant DT, with an accumulator variable that persists across frames. This means that if there is not enough time to do a complete Tick, the remaining time is added to the next frame and an extra Tick could be carried out then. Essentially this means that a tick only happens once a certain fixed amount of time has passed. This allows us to manage the operating load of the physics engine more than the previous example while still maintaining an accurate time line(T).

3.1.4 Game Loop Independant

The previous three methods are attempts to decouple the physics simulation from the framerate of the running application. It may have occurred to you already that this can be done in a much more substantial way: Multithreading, i.e keeping the physics simulation running in the background. This would be more of a concurrency task than physics, so it is not covered in this module.

3.1.5 Tutorial Exercises

Exercises

1. Complete the 3 time-step examples (timing_simple.cpp, unfixed_timestep.cpp, unfixed_timestep.cpp)
2. Experiment with limiting the frame-rate and doWork() delay, how does this effect the physics simulation?

Advanced Exercises

- Implement the physics loop in a separate cpu thread.

3.2 Integration

Now that we have some of the more technical framework code covered for instigating a physics Tick, we will now look at what actually happens in the physics world during a tick.

Keep in mind that the tick function takes the current time T , and a delta time DT . What we are asking the physics engine to do is calculate where everything will be at time $T+DT$.

To do this, all the general rules of Newtonian and others are applied to the current state of the world. Unfortunately, the fact that we are running ticks at discrete points in time can lead to inaccurate results when run over many Ticks. To combat this, different types of *Integration* methods are used.

This is the same theory as integration in the field of calculus, as we are attempting to solve ordinary differential equations in its most basic forms. We are going to hyper-warp over the mathematical proofs that define most of this and stick to two simple concepts: The concept of Order (First order, Second Order...), and the concept of Error(Local and Global).

Let's look at the most simplest form of Integration to begin with before we get really stuck in

3.2.1 The Euler method

```
1 void Tick(const double t, const double dt) {
2     for (each object in scene)
3     {
4         vec3 acc = calculateAcceleration(object);
5         object.velocity += acc * dt;
6         object.position += object.velocity * dt;
7     }
8 }
```

3.2.2 The Verlet method

Verlet is an integration method that was created for use in the Atomic and Molecular Dynamics fields. It is based on a set of mathematical foundations, principally Taylor Expansion. Fortunately for us, the resulting algorithm is only a short step away from Euler, the short explanation is that we no longer store the velocity of objects, instead we keep track of the previous and current position, and use this to calculate the velocity each Tick.

```
1 void Tick(const double t, const double dt) {
2     for (each object in scene)
3     {
4         vec3 acc = calculateAcceleration(object);
5         vec3 velocity = object.position - object.position_prev;
6         object.position_prev = object.position;
7         object.position += velocity + acc * dt * dt;
8     }
9 }
```

This small change actually makes this a second order function, which is much better at conserving internal Energy. Verlet can make solving constraints trivially, as we now do not have to modify velocities, only positions. The downside is that it requires a significantly smaller timestep to settle on solutions.

3.2.3 The Runge-Kutta (RK4) method

Technically all the integrations mentioned prior to this are versions of Runge–Kutta methods, however the RK4 integrator is usually what is being referred to when Runge-Kutta is mentioned. This is a Fourth order technique and can therefore deal with drastic changes in acceleration more accurately. In laymen's terms, RK4 calculates 4 positions(P) and Velocity(V) spread equally across the timespan of Δt . It then sums the results together with a higher weight applied to the middle values than the edges.

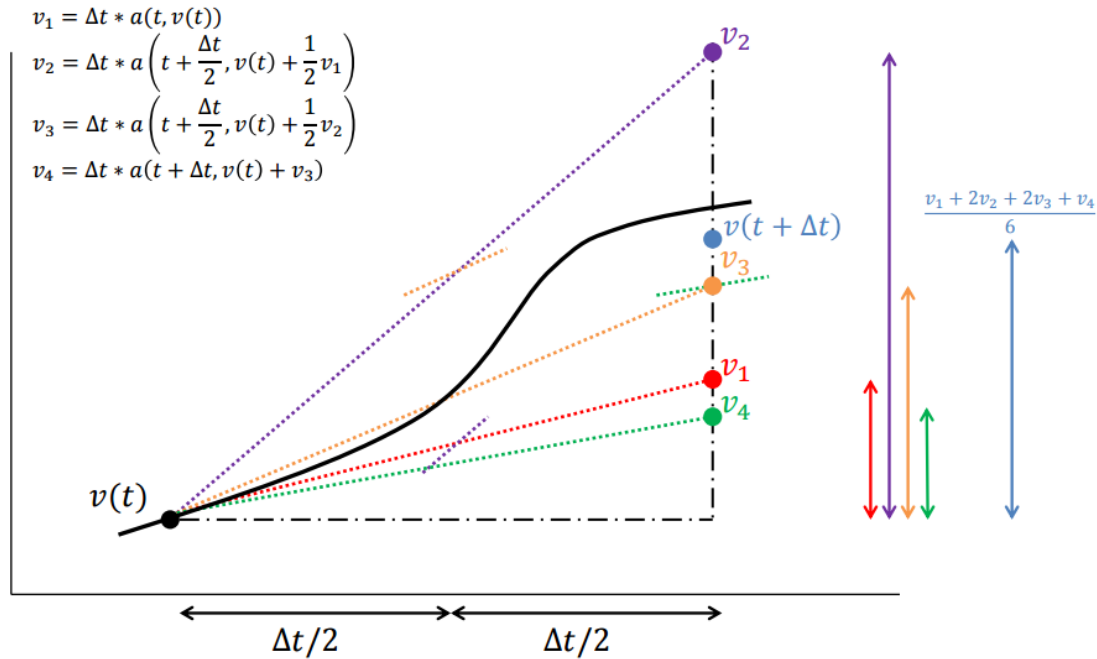


Figure 3.1: **RK4 steps compared**[4] - Black is the ground truth, blue is the RK4 result

```

1  p1 = position;
2  v1 = velocity;
3  a1 = acceleration( p1, v1);
4  //Calculate P and V, at half dt
5  p2 = p1 + v1 * dt / 2;
6  v2 = v1 + a1 * dt / 2;
7  //Calculate P and V, at half dt, with new V and A
8  a2 = acceleration( p2, v2);
9  p3 = p1 + v2 * dt / 2;
10 v3 = v1 + a2 * dt / 2;
11 //Calculate P and V, at full dt, with new V and A
12 a3 = acceleration( p3, v3);
13 p4 = p1 + v3 * dt;
14 v4 = v1 + a3 * dt;
15 a4 = acceleration( p4, v4);
16 //now we have 4 counts for P and V accross Dt, now "merge" down to one
17 position += (v1 + 2 * v2 + 2 * v3 + v4) * dt / 6;
18 velocity += (a1 + 2 * a2 + 2 * a3 + a4) * dt / 6;

```

Listing 3.1: RK4 Pseudocode

3.2.4 Tutorial Exercises

Exercises

1. Complete the 3 integration examples (Euler.cpp, Verlet.cpp, RK4.cpp)
2. Add additional acceleration functions in addition to Gravity

Advanced Exercises

- Implement a method for measuring the accuracy of each integration method.

Chapter 4

Particles

Chapter 5

Collision Detection

Chapter 6

Rigid Body

Chapter 7

Appendix

7.1 Bibliography

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