Profile Laboratory Notes Chris Gregory and Deanna Bessy

#### Version 0

Inital UM Emulator after project 6

Midmark.um

# Instructions: 101,046,726,745

User time: 18.69s

• Sandmark.um

User time: 745.60s (~12min)

### **Version 1**

This step did not significantly improve the performance. Compiler optimization switched to level 2

Midmark.um

# Instructions: 101,046,726,745

User time: 18.69s

User time/initial user time: 1 User time/previous user time: 1

• Sandmark.um

User time: 710.96s (~12min)

User time/initial user time: 0.9535 User time/previous user time: 0.9535

# Version 2

Bottleneck was the lookup of Atoms as IDs in Hanson tables and the indirection of pointers in the Hanson sequences. We corrected this by replacing tables and sequences with arrays.

Midmark.um

# Instructions: 18,240,484,439

User time: 1.71s

User time/initial user time: 0.0915 User time/previous user time: 0.0915

• Sandmark.um

User time: 37.92s

User time/initial user time: 0.0508 User time/previous user time: 0.0533 Bottleneck was the speed of accessing functions from other modules. We corrected this by putting all of the functions from all modules in one file.

### • Midmark.um

# Instructions: 5,348,912,432

User time: 1.61s

User time/initial user time: 0.0861 User time/previous user time: 0.9415

# • Sandmark.um

User time: 34.17s

User time/initial user time: 0.0458 User time/previous user time: 0.9011

# Version 4

Bottleneck was the bitpack functions from the included bitpack module. We corrected this by moving some whole bitpack functions into our file and using some code from other bitpack functions that did bit shifting.

#### • Midmark.um

# Instructions: 5,337,276,241

User time: 0.64

User time/initial user time: 0.0342 User time/previous user time: 0.3975

# • Sandmark.um

User time: 9.95s

User time/initial user time: 0.0133 User time/previous user time: 0.2911