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# Computational Evolution & Digital Organisms

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A look at a subset of Artificial Life

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# Computational Evolution

- Attempts to elucidate principles of evolution
    - Builds models of self-replicating organisms
      - Computational cost limits physical fidelity of the model.
      - Digital or chemical models
    - Mutation creates variation in populations
    - Reproduction can be sexual or asexual
    - Ability to (out) reproduce its genome is the usual fitness measure
      - For some research, other fitness measures are used.
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# Not to be Confused With Evolutionary Computing

- A Search Technique inspired by biology
    - Points in search space represented as “genomes”
    - Crossover produces new points in search space
    - Mutation ensures variety
      - Ensures more of search space is sampled
    - Fitness function determines which subset of population become progenitors
    - Larger populations increase coverage of space.
    - Search usually walks through “invalid” points
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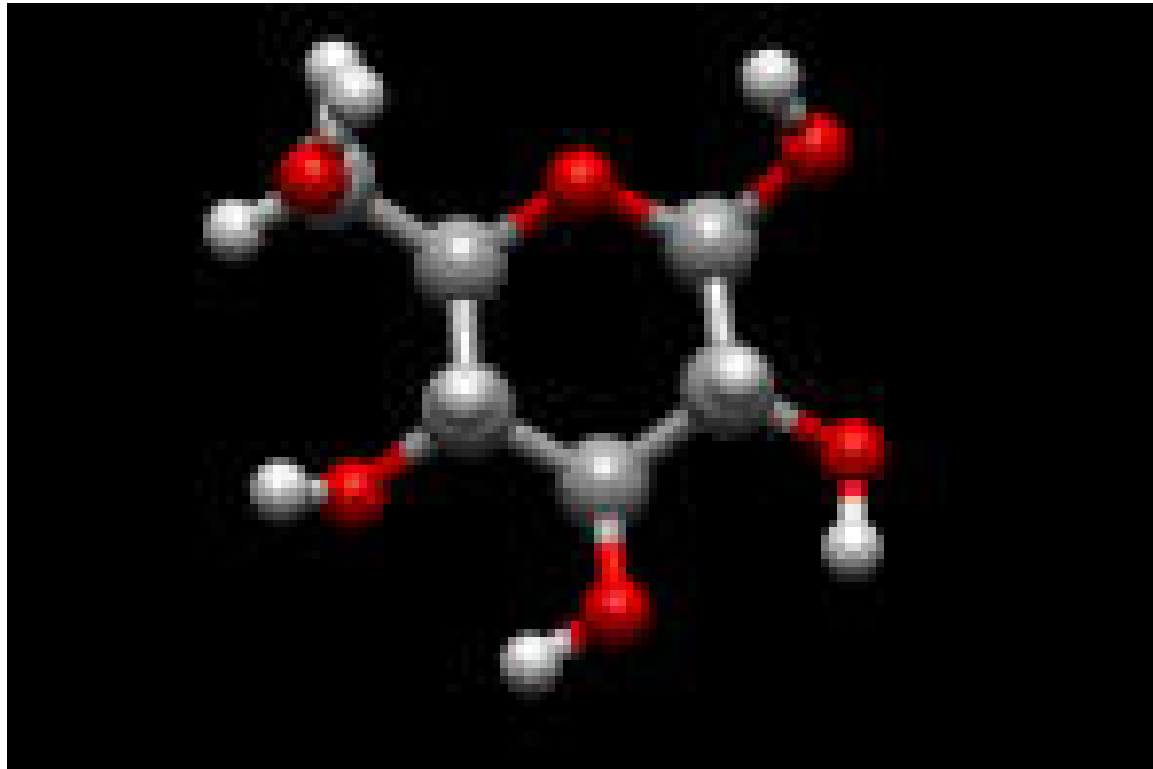
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# Overview of Talk

- Motivation: The complexity of cellular life
  - Tierra and the evolution of digital organisms
  - Avida and other Tierra inspired work
  - Lessons/Future Research
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# Complexity of Cellular Life I: Metabolism of Glucose to produce ATP



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# Molecules of glucose pathway.

- PDB Molecule of the Month The Glycolytic Enzymes
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## It's all chicken and egg

- Where did glucose come from?
  - Where did all those intermediate products come from?
  - Where did all those wonderful enzymes come from?
  - Take away any of the enzymes, and the system collapses.
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# Complexity of Regulatory Mechanisms

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# Nature made this from

- Molecules with differential binding affinities for DNA.
  - Overlapping control regions.
  - Positive and negative feedback.
  - Cooperative binding.
  - How did it make the recipe?
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# Tierra, a Platform for Digital Evolution

- Design Requirements/Inventions:
    - Organisms must be self-reproductive
    - Ability to out-reproduce the competition only fitness criteria
      - Avoids “artificial” fitness functions.
    - Control (jumps/calls) is effected through *templates* and *targets*, which are complementary “bit strings”
      - Jump nop1 nop0 nop1 goes to nop0 nop1 nop0
    - Organisms sense the environment
      - Dynamic “fitness” function
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# Tierra's Digital Organisms

- Each organism (cpu) has
    - 4 registers (A, B, C, D)
    - Instruction pointer
    - 10 word stack
  - Time slicing “implements” parallel organisms
  - When space for new organisms is needed, the oldest organisms are reaped (as a rule).
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# Tierra's Instruction Set

## ■ Data Movement

- PushA, PopA, PushB, PopB, etc for C and D
- MOVDC ( $D \leftarrow C$ ), MOVBA, COPY ([A] to [B])

## ■ Control

- JumpO, JumpB, Call, Ret, IfZ, nop0, nop1

## ■ Calculation

- subcab, subaac, inca, incb, decc, incd, zero, not, shl

## ■ Biological and Sensing

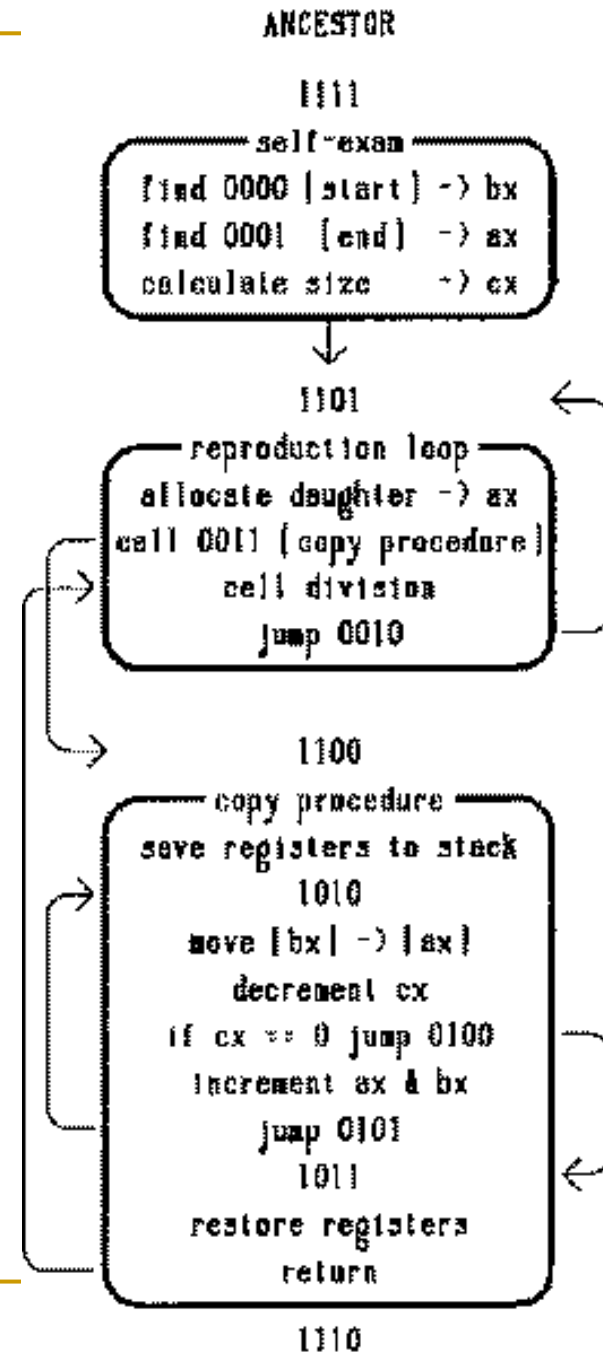
- adr, adrb, adrF, mal (allocate memory), divide
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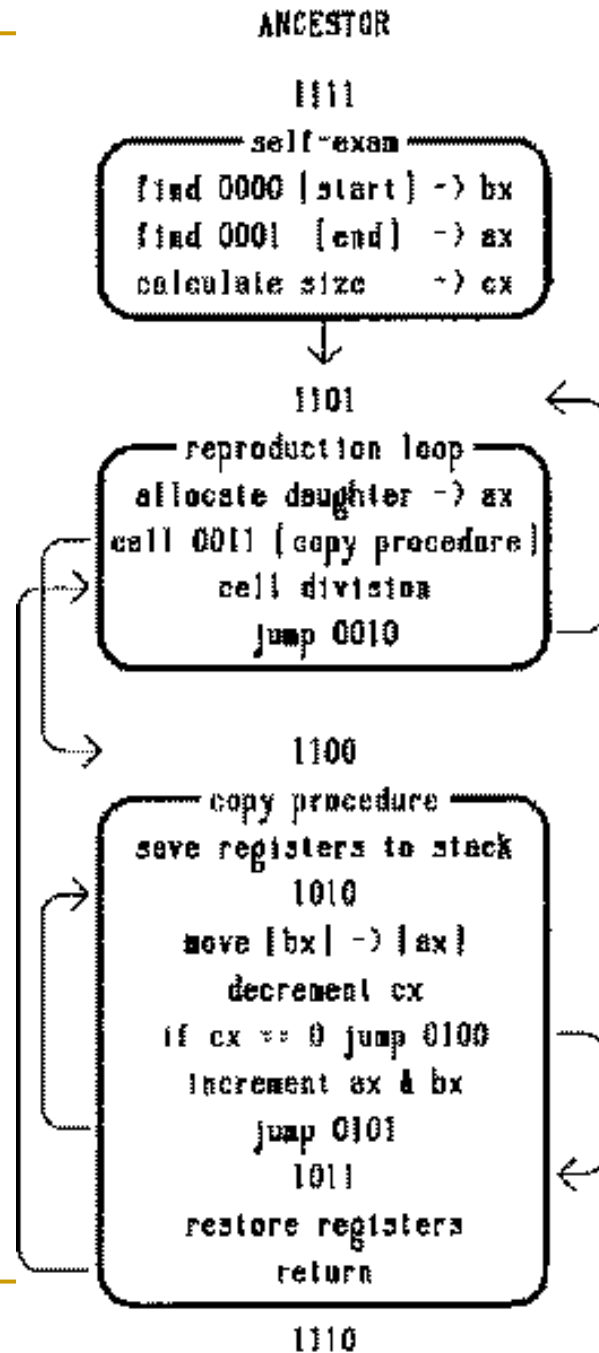
# Mutational Sources

- A copy error every  $X$  copy instructions
  - Cosmic rays
    - A bit in the soup gets flipped every  $Y$  instructions
    - Works because no cells are autosomes
    - Biased, not random
  - Probabilistic results of instructions
    - Every so often an instruction misfires
    - E.g., `incA` adds 2
  - No Insertion/deletions
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# The Tierran Ancestor



# The Tierran Ancestor



Lots of redundancy

- Labels can be shortened
- Different control constructs
- Cells only replicate once or twice
- Templates can be labels
- Various return addresses can be used
- Control can use any matching code

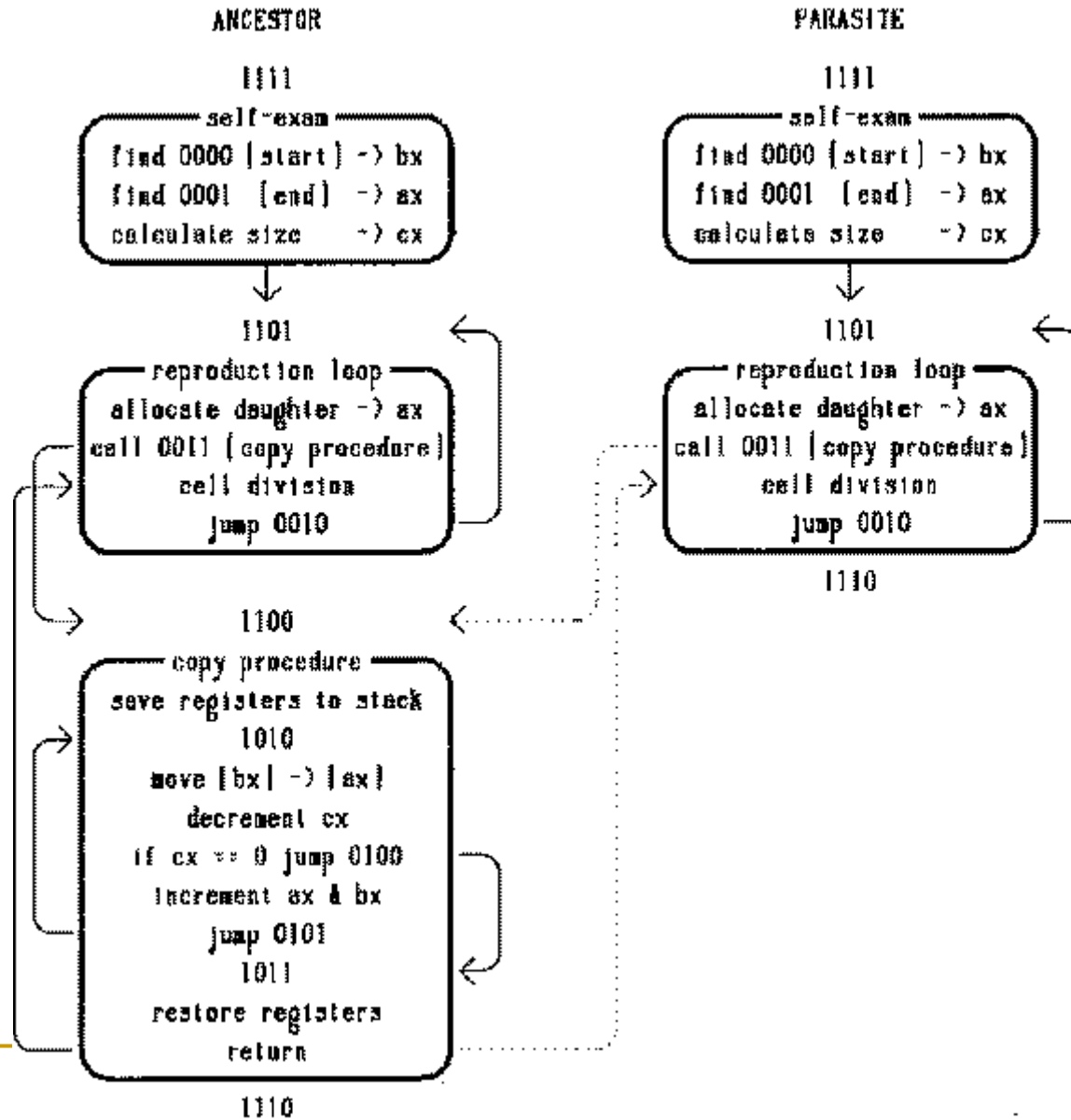
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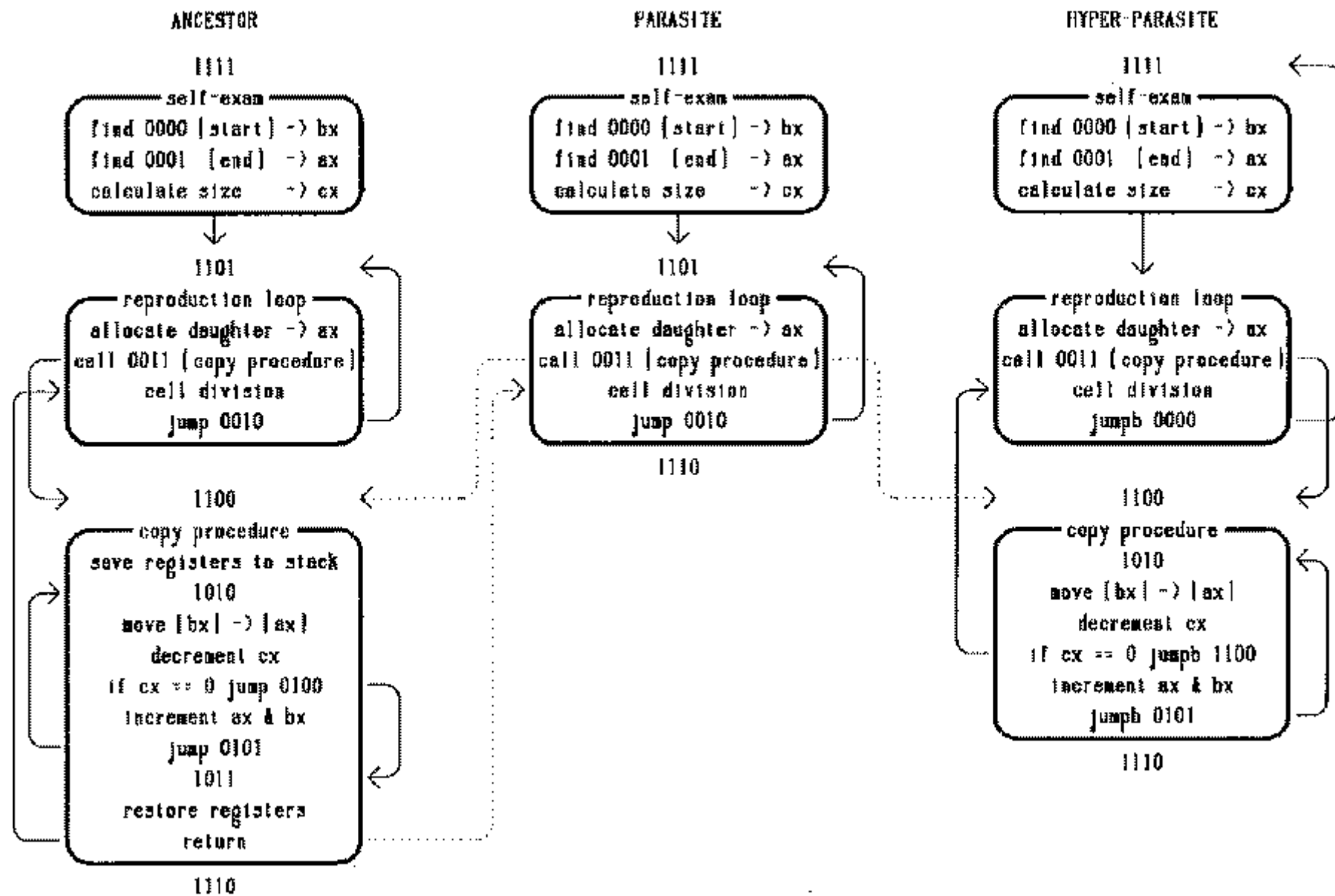
# Ancestor Code

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# Evolution in Action: Parasites!!





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## An interesting chicken-and-egg mutation

- <C = size, B=@self>
  - nop1 nop1 nop0 nop1
  - mal
  - call nop0 nop0 nop1 nop1
  - divide
  - jump nop0 nop0 nop1 nop0
  - ifz
  - nop1 nop1 nop0 nop0
  - <copy loop>
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## An interesting chicken-and-egg mutation

- <C = size, B=@self>
  - nop1 nop1 nop0 nop1
  - mal
  - call nop0 nop0 nop1 nop1
  - divide
  - pushb (was jump) nop0 nop0 nop1 nop0
  - ifz
  - ret (was nop1) nop1 nop0 nop0
  - <copy loop>
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# A Copy-Once Parasite

- Stays just ahead of the reaper
    - ❑ `nop1 nop1 zero not0 shl shl movdc`
    - ❑ `adrb nop0 nop0 pushc nop0`
    - ❑ `subaac`
    - ❑ `movba pushd nop0`
    - ❑ `adr nop0 nop1`
    - ❑ `inca`
    - ❑ `subcab pusha nop1 pushd nop1`
    - ❑ `mal`
    - ❑ `call nop0 nop0 nop1 nop0`
    - ❑ `divide`
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# Two chances to find a copy loop

- ❑ <C = size, B = @self>
  - ❑ mal pusha call movii pusha
  - ❑ call nop0 nop0 nop1 nop1
  - ❑ divide movii
  - ❑ pusha
  - ❑ mal
  - ❑ call nop0 nop0 nop1 nop1
  - ❑ divide mal subaac nop1
  - ❑ ret zero nop1 zero (jumps to start of daughter)
  - ❑ nop1 nop1 nop1 nop0
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SYMBIONTS

1111

```

self-exam
find 0000 (start) -> bx
find 0001 (end) -> ax
calculate size -> cx
jump 0010
    
```

1111

```

self-exam
find 0000 (start) -> bx
find 0001 (end) -> ax
calculate size -> cx
jump 0010
    
```

1100

```

copy procedure
save registers to stack
move [bx] -> [ax]
decrement cx
if cx == 0 jump 0100
increment ax & bx
jumpb 0101
1011
restore registers
return
    
```

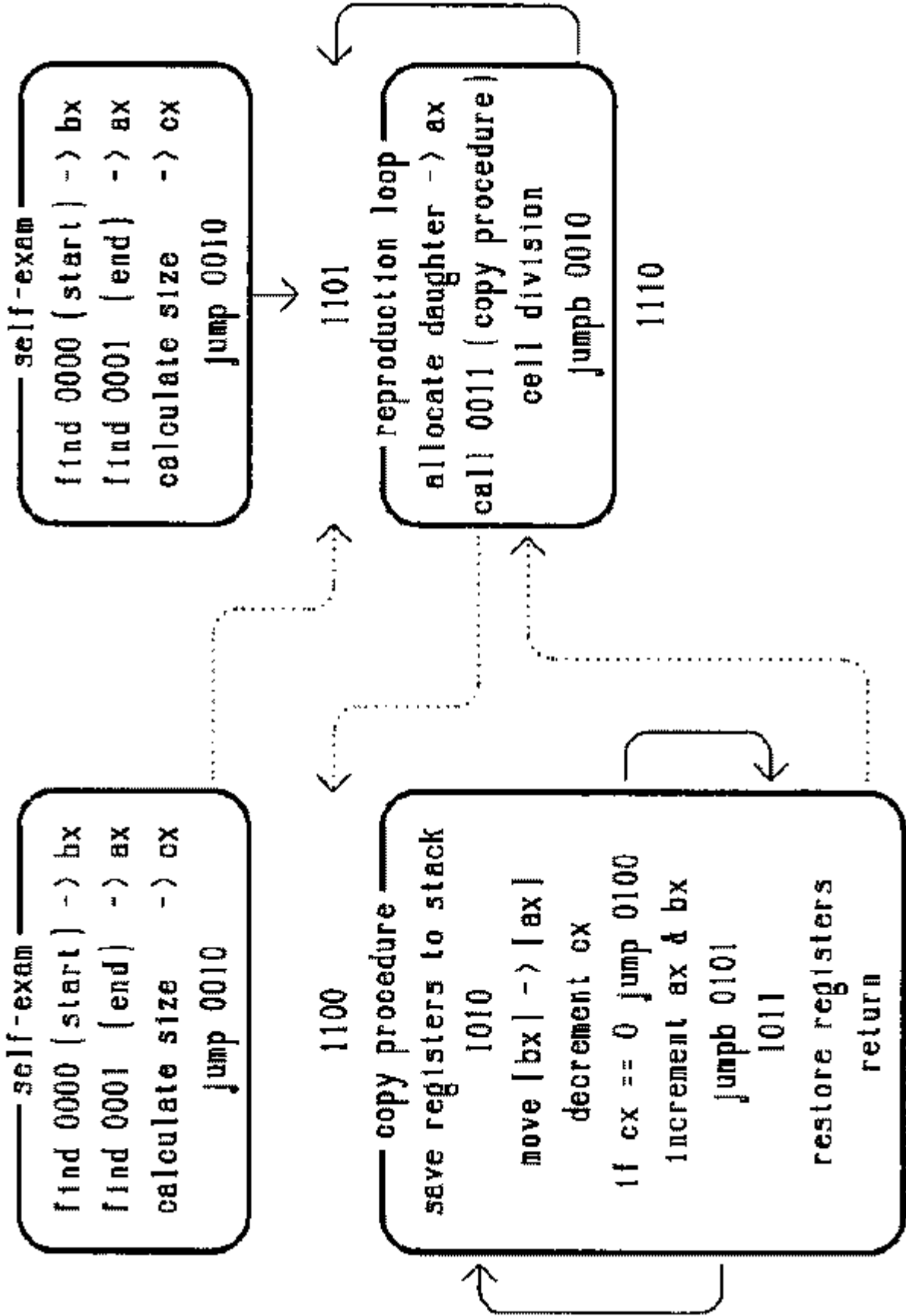
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```

reproduction loop
allocate daughter -> ax
call 0011 (copy procedure)
cell division
jumpb 0010
    
```

1110

1110



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## Feature or Bug?

### CPU is independent of genome

- A very small self-replicating parasite (15 long)
    - Nop1
    - Adrb nop0
    - MovBA
    - Adrf nop0 nop0
    - subAAC
    - Jump nop0 nop0 nop1 nop0
    - Nop1 nop1
  - Even smaller viable program:
    - Nop1
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# Feature or Bug?

## Non-local effects

- A template can match *any* nearby target
  - A request for memory can kill *any* organism, even one “fitter”
  - A daughter cell can be placed *anywhere*
  - Allocating a large amount of memory for a daughter can kill tens of organisms, creating a dieoff
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Feature or Bug?

Spaghetti Code is a Frequent Occurrence

- Symbionts arise quite frequently
- When a target is mutated, the target in another cell is used.

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## Bug or Feature?

### Parasites require necrophilia

- Instructions are left in memory when an organism is reaped.
- “Parasites” keep using these instructions.

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## Bug or Feature?

### Sloppy replicators instead of Indels

- Teirra lacks insertion/deletion mutations
    - Biology uses indels
    - Harder to remove instructions without deletions
    - Harder to make room for new instructions
  - Tierra makes up for it with sloppy replicators that move instructions around willy nilly
    - Buy maybe this is needed anyway?
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# Is Sloppiness needed to Bootstrap Complexity?

- Sloppiness (ad-hoc) mixing gave us
    - Mitochondria (ingestion without digestion)
    - Chloroplasts in bacteria (same story)
    - Gene mixing (via viruses)
    - Diploidy from Haploidy
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# Avida

- Inspired by Tierra, but
    - Controlled instruction pointers (less slopiness)
    - Insertion/Deletion mutations
    - 2 dimensional grid of organisms, not instructions
    - Only local next-neighbor effects
    - Fitness functions to augment reproduction
  - Experiments to test biological theories
    - Evolution of Complexity
    - Evolution of Complex Functions
    - Relationship among evolution rate and landscape
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# Digital Biosphere

- Inspired by Tierra/Avida but
    - Focus is on evolutionary trajectories.
      - Are there principles regarding these trajectories?
    - Will exploit the constraints of physics
      - Conservation Laws!
      - Energy requirements and metabolism
    - Will eventually move to chemical modeling to get closer to biology.
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# Lessons

- Evolution finds corners of the search space
    - If you build it, they will exploit it
    - Complexity comes from exploiting environment
  - Co-evolution makes the problem interesting and different
    - Changing fitness functions
  - Designing a system for open-ended evolution is still very much an open-ended problem.
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What's it all mean?

We have a source of new insights

- Watching evolving dynamical systems give insight and ideas.
  - Biologists aren't trained to do this.
  - Many insights will be gained that will eventually transfer over to biological thinking
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# Last Thought

- Is the complexity of the phage lambda lyse/lytic growth mechanism any more or less complex than the programs that Tierra was evolving?
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