IBRI Colloquium, 22 Jan 1991 Dr. Robert C. Newman Biblical Theological Seminary

# COMPUTER SIMULATIONS OF EVOLUTION

## Introduction

Not a literature search Not covering origin of life question tho 2 programs on diskette are self-reproducing automata REPRO - Langton's automaton in my *JASA* Spr 88 paper BYL - Byl's in his *JASA* Spr 89 paper both these can be "mutated" by manipulating data for structure & transition rules Not dealing with competition & spread of varieties good deal has been done on ecology/population genetics Rather, a description & investigation of three programs that relate to the **mechanism** of evolution: -- two described by Richard Dawkins in his *Blind Watchmaker* (1986), 46-75 -- one devised by self these 3 programs also on diskette available from IBRI for \$5

## **Program BIOMORPH**

Describe:

program, slightly simplified from Dawkins, for building "organisms" from genetic information, selecting among mutants

gene is sequence of eight small integers

generates "tree" controlling branch length, angles,

# of levels of branching, with mirror symmetry

given original gene, program constructs all "one-step" mutations, displays on screen operator selects which mutant to succeed parent

### Lessons from BIOMORPH:

shows how:

mutation operates on DNA

selection operates on developed form, not on DNA

see that:

identical forms can conceal diff genetics leaving room for neutral mutation

## **Program SHAKES**

Describe:

Dawkins seeking to circumvent "monkeys typing Shakespeare" problem of enormous times involved

choose target sentence/phrase

start with gibberish of same length

mutate gibberish, selecting mutant/parent which is closer to target to be new parent gibberish converges to target much faster than if monkeys were typing randomly

Dawkins gets convergence in typically 40-70 generations Dawkins' version: Not described in detail, so can't tell how he generated mutants, how many mutations per generation My version: One mutant each generation, compared w/ parent Better of mutant/parent survives I get much slower convergence, taking over 1000 generations Lessons from SHAKES. shows that a "rachet mechanism" does work important reason why many convinced evolution must be correct but this is "guided evolution," which is considerably more efficient than even artificial selection, to say nothing of natural selection! does not solve time question which version is more realistic? mutation rate in eukaryotes is  $10^{-8}$  per replication both ignore time involved for mutant to take over population! my version suggests a problem for mutating into complex or optimal structures: last pieces of puzzle are highly constrained **Program MUNSEL** Describe: simulate mutation & natural selection by analogy with human language letter string is both gene and organism mutation is random change in content and/or length selection is "naturalized" by requiring that each grouping in string be an English word current version has operator do selecting, but comparing with a spell-checker would be more objective generates words of 1-4 letters rather easily relative frequency of space character (and nature of selection) tends to keep words short little success in getting intelligibility in 100s of steps

Lessons from MUNSEL:

complex organisms involve hierarchies of structure

somewhat like intelligible writing

letters > words > phrases/sentences > paragraphs

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mutation only works at lowest level
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nucleotides <=> letters

so becomes tougher to get anything acceptable as we move up hierarchy

non-selected mutation => gibberish
neutral mutations spread only by random walk
functional isolation seen here (as in terrain analogy)
many combinations cannot be reached by single mutations from acceptable
smaller groups
what is the relative size of islands of intelligibility vs oceans of gibberish
for each level of hierarchy?
can you really get there from here?
complex organs/organisms
crossing higher levels of bio classification