ITERATIONS, LOOPS, RECURSIONS, AND GENETIC PROGRAMMING PROBLEM SOLVER (GPPS)

ITERATION

- Four parts
 - initialization branch
 - termination branch
 - body of iteration (work-performing branch)
 - update branch
- Difficulties
 - nested iterations
 - unsatisfiable termination predicates
- Rationing

ITERATION

DIFFERENT APPROACHES

- DU ("Do Until") operation (GP-1)
 - open-ended number of steps in an iteration
 - open-ended number of iterations
- Automatically defined iterations (ADIs) (GP-2). Also called "restricted iteration"
 - There is only has a iteration-performing branch (IPB)
 - Iteration is over a known, fixed set
 - protein or DNA sequence (of varying length)
 - time-series data
 - two-dimensional array of pixels
- Automatically defined loops (ADLs) (GP-3)
 - loop initialization branch
 - loop termination branch
 - loop body branch
 - loop update branch

AUTOMATICALLY DEFINED ITERATIONS (ADIs)

- Overall program consisting of an automatically defined function ADFO, an iteration-performing branch IPBO, and a result-producing branch RPBO.
- Iteration is over a known, fixed set
 - protein or DNA sequence (of varying length
 - time-series data
 - two-dimensional array of pixels

TRANSMEMBRANE SEGMENT IDENTIFICATION PROBLEM

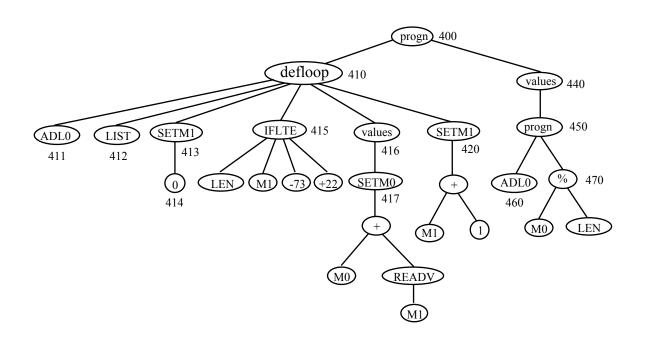
- Goal is to classify a given protein segment as being a transmembrane domain or non-transmembrane area of the protein
- Generation 20 Run 3 Subset-creating version
 - in-sample correlation of 0.976
 - out-of-sample correlation of 0.968
 - out-of-sample error rate 1.6%

```
(progn (defun ADFO ()
(values
(defun ADF1 ()
(ORN (ORN (I?) (H?)) (ORN (P?) (G?))) (ORN (ORN
(ORN (Y?) (N?)) (ORN (T?) (Q?))) (ORN (A?) (H?)))))
(defun ADF1 ()
(values (ORN (ORN (ORN (A?) (I?)) (ORN (L?) (W?)))
(ORN (ORN (T?) (L?)) (ORN (T?) (W?)))))
(defun ADF2 ()
(values (ORN (ORN (ORN (ORN (D?) (E?)) (ORN (ORN
(ORN (D?) (E?)) (ORN (ORN (T?) (W?)) (ORN (Q?)
(D?)))) (ORN (K?) (P?)))) (ORN (K?) (P?))) (ORN (T?)
(W?))) (ORN (ORN (E?) (A?)) (ORN (N?) (R?)))))
(progn (loop-over-residues
       (SETMO (+ (- (ADF1) (ADF2)) (SETM3 MO))))
(values (% (% M3 M0) (% (% (% (- L -0.53) (* M0 M0))
(+ (% (% M3 M0) (% (+ M0 M3) (% M1 M2))) M2)) (% M3
MO))))))
```

COMPARISON OF EIGHT METHODS FOR SOLVING TRANSMEMBRANE SEGMENT IDENTIFICATION PROBLEM

Method	Error
von Heijne 1992	2.8%
Engelman, Steitz, and Goldman 1986	2.7%
Kyte and Doolittle 1982	2.5%
Weiss, Cohen, and Indurkhya 1993	2.5%
GP + Set-creating ADFs (GP-2)	1.6%
GP + Arithmetic-performing ADFs (GP-2)	1.6%
GP + ADFs + architecture-altering	1.6%
operations for subroutines (GP-3, chapter 16)	
GP + ADFs + architecture-altering	1.6%
operations for both subroutines and	
iterations (GP-3, chapter 17)	

AUTOMATICALLY DEFINED LOOPS EXAMPLE OF A PROGRAM WITH A FOUR-BRANCH AUTOMATICALLY DEFINED LOOP (ADL0) AND A RESULTPRODUCING BRANCH

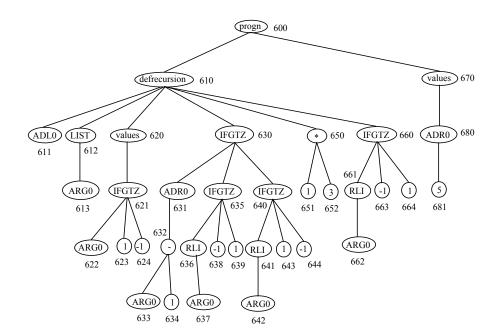


RECURSION

- Recursion is difficult for GP because
 - Small changes in the STRUCTURE of a recursive program often lead to vastly different PERFORMANCE of the program and therefore very different values of fitness
 - Small changes in base case (initial value) as in logistic function
 - Small changes in transition rule
- Practical problems of time (like iteration) because of potentially
 - Infinite recursion
 - Near-infinite recursion

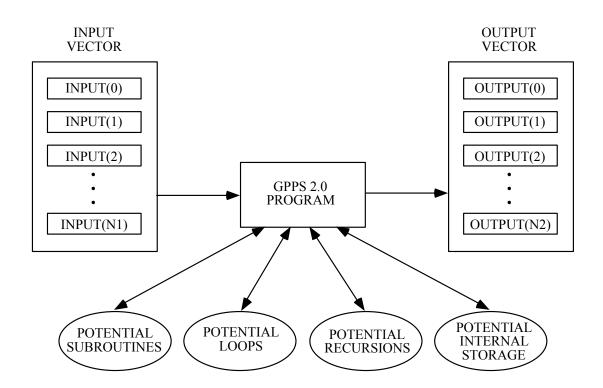
AUTOMATICALLY DEFINED RECURSION (ADR0) AND A RESULT-PRODUCING BRANCH

- recursion condition branch, RCB
- recursion body branch, RBB
- recursion update branch, RUB
- recursion ground branch, RGB



GENETIC PROGRAMMING PROBLEM SOLVER (GPPS)

— VERSION 2.0



GENETIC PROGRAMMING PROBLEM SOLVER (GPPS 2.0)

Objective:	Create, using the Genetic
	Programming Problem
	Solver 2.0, a computer
	program that takes the
	values of one independent
	floating-point variable x in
	the input vector and deposits
	a value into the output vector
	that closely matches the
	quadratic polynomial
	$2.718x^2 + 3.1416x$.
Program architecture:	One result-producing
	branch, RPB. Automatically
	defined loops, automatically
	defined recursions,
	automatically defined stores,
	and automatically defined
	function(s) and their
	arguments will be created
	during the run by the
	architecture-altering
	operations.
	$F_{\text{rpb-initial}} = \{+, -, *, \%, \}$
result-producing branches:	IFLTE, TOR, TAND, TNOT,
	RLI, WLO, FLOOR}.

Initial terminal set for the	
result-producing branches:	NINPUTS, NOUTPUTS,
	INDEX}.
Initial function set for the	No automatically defined
	functions in generation 0.
functions:	$F_{adf-initial} = \phi$.
Initial terminal set for the	·
	functions in generation 0.
	$T_{adf-initial} = \phi$.
	No automatically defined
	loops in generation 0. F _{adl-}
automaticany ucinicu loops.	
	$initial = \phi$.
	No automatically defined
automatically defined loops:	loops in generation 0. Tadl-
	$_{initial} = \phi$.
Initial function set for	No automatically defined
	recursions in generation 0.
recursions:	$F_{adr-initial} = \phi$.
Initial terminal set for	No automatically defined
	recursions in generation 0.
recursions:	$F_{adr-initial} = \phi$.
	•
Potential function set for the	Po booming /
result-producing branches:	SWB0, SWB1, ADF0, ADF1,
	ADF2, ADF3}.
Potential terminal set for the	$T_{\text{rpb-potential}} = \{LBB0, SRB0, $
result-producing branches:	SRB1}.
Potential function set for the	$ F_{adf-potential} = \{ADF0, ADF1, $
automatically defined	ADF2, ADF3}.
functions:	
ı	

Potential terminal set for the	$T_{adf-potential} = \{ARG0, ARG1,$
automatically defined	_
functions:	INDEX, ←bigger-reals}.
Potential function set for	
automatically defined loops:	ADL2, ADL3}
Potential terminal set for	T _{adl-potential} = {NINPUTS,
automatically defined loops:	NOUTPUTS, INDEX, ←bigger-
	reals}.
Potential function set for	$F_{adr-potential} = \{ADR0, ADR0,$
automatically defined	ADR1, ADR2, ADR3}.
recursions:	
Potential terminal set for	$ T_{adr-potential} = \{NINPUTS, $
automatically defined	NOUTPUTS, INDEX, ←bigger-
recursions:	reals}.
Fitness cases:	20 randomly chosen values
	of the independent variable x
	between -1.0 and +1.0.
Raw fitness:	Raw fitness is the sum, over
	the 20 fitness cases, of the
	absolute value of the
	difference between the value
	deposited in the output
	vector and the value of the
	quadratic polynomial
	$2.718x^2 + 3.1416x$.
Standardized fitness:	Same as raw fitness

Hits:	The number of fitness cases (0 to 20) for which the value deposited in the output vector is within 0.01 of the value of the quadratic polynomial 2.718x ² + 3.1416x.
Wrapper:	None.
Parameters:	$M=120,000.$ $G=1,001.$ $NINPUTS=1.$ $NOUTPUTS=1.$ $Q=2,000.$ $D=60.$ $B=2\%.$ $N_{rpb}=1.$ $S_{rpb}=500.$ $S_{adf}=100.$ $N_{max-adf}=4.$ $N_{max-argument-adf}=2.$ $N_{min-adf-arg}=0.$ $N_{max-adl}=1.$ $S_{adl}=100.$ $N_{max-argument-adl}=0.$ $N_{min-argument-adl}=0.$ $N_{max-adl-executions}=3.$ $N_{max-adr-adl-executions}=1.$ $S_{adr}=100.$ $N_{max-argument-adr}=0.$ $N_{max-argument-adr}=0.$ $N_{max-adr-executions}=9.$ $N_{max-adr-executions}=9.$ $N_{max-ads}=2.$
Result designation:	Best-so-far pace-setting individual.
Success predicate:	A program scores 20 hits.