

Evolutionary Identification of Wind Turbine Dynamics

William La Cava

Advisors:

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Lee Spector²

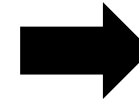
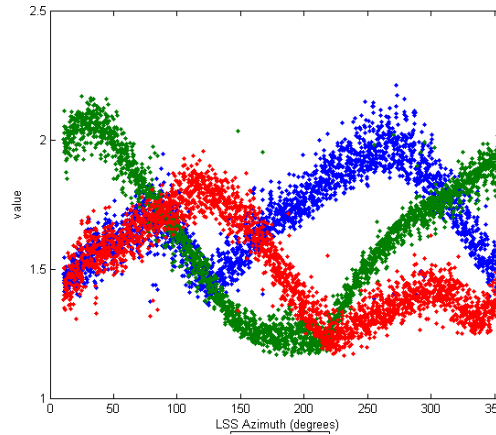
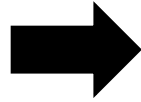
Matthew Lackner¹

¹*M&IE, UMass Amherst*

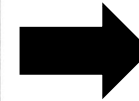
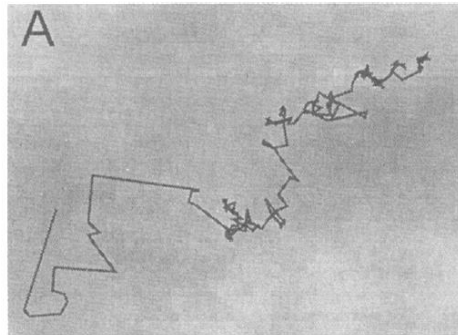
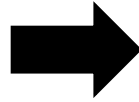
²*Cognitive Science, Hampshire College*



System Identification



$$y = f(t, x, a, b, c)$$



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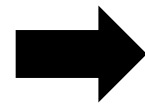
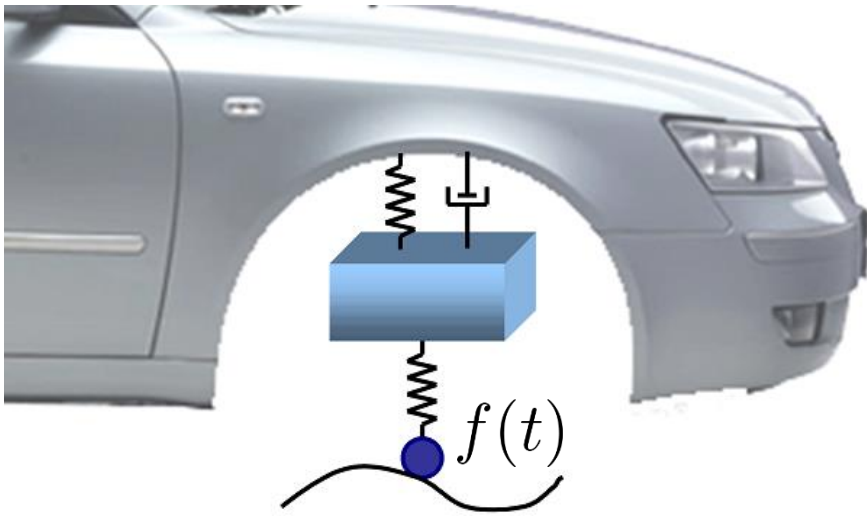
System Identification

First Principles

$$\sum F = m\ddot{x}$$

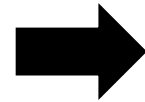
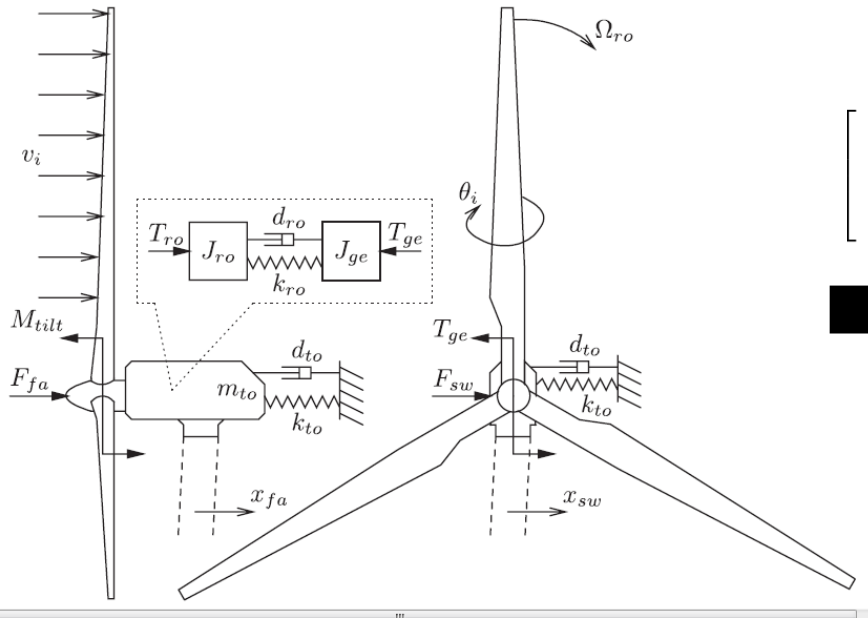


System Identification



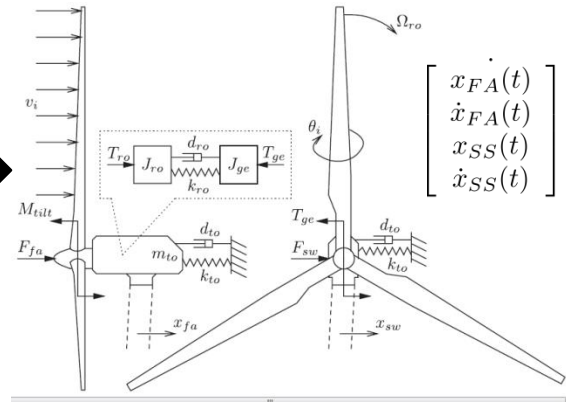
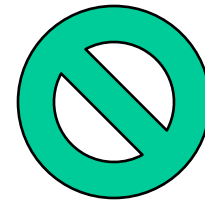
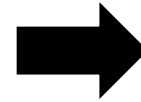
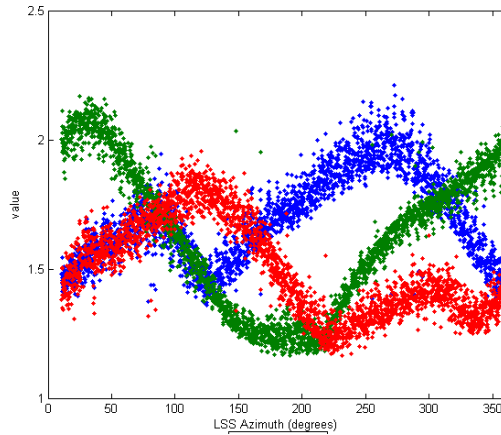
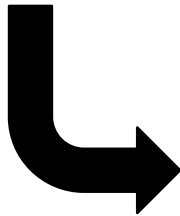
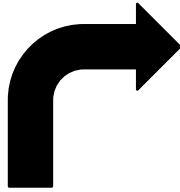
$$m\ddot{x} + c\dot{x} + kx = f(t)$$

System Identification



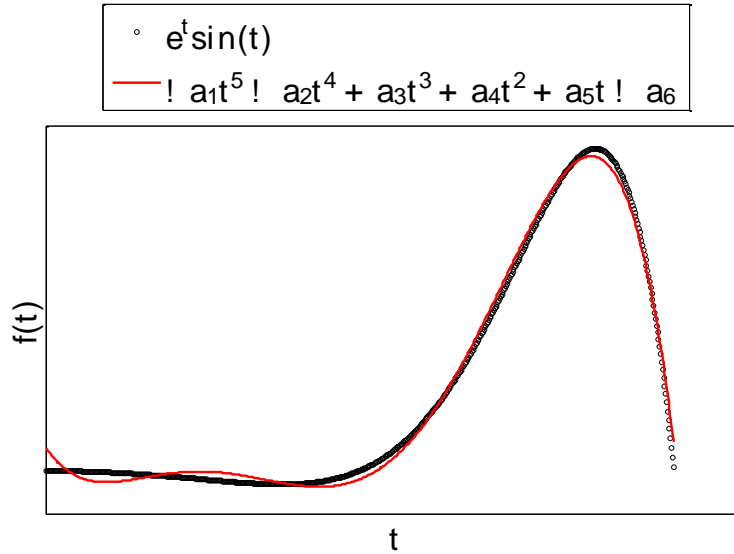
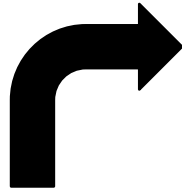
$$\begin{bmatrix} \dot{x}_{FA}(t) \\ \dot{x}_{FA}(t) \\ \dot{x}_{SS}(t) \\ \dot{x}_{SS}(t) \end{bmatrix} = \begin{bmatrix} 0 & 1 & 0 & 0 \\ -\frac{k_{to}}{m_{to}} & \frac{81R}{32H^2} \frac{h_{Mz}}{m_{to}} - \frac{d_{to}}{m_{to}} - \frac{3h_{Fx}}{m_{to}} & 0 & 0 \\ 0 & 0 & 0 & 0 \\ 0 & -\frac{27R}{16H^2} \frac{h_{Fz}}{m_{to}} & -\frac{k_{to}}{m_{to}} & -\frac{d_{to}}{m_{to}} \end{bmatrix} \begin{bmatrix} x_{FA}(t) \\ \dot{x}_{FA}(t) \\ x_{SS}(t) \\ \dot{x}_{SS}(t) \end{bmatrix} + \begin{bmatrix} 0 & 0 \\ 0 & \frac{3k_{Fx}}{m_{to}} \\ 0 & 0 \\ \frac{3}{2H} \frac{1}{m_{to}} & 0 \end{bmatrix} \begin{bmatrix} T_{gen} \\ \beta \end{bmatrix} + \begin{bmatrix} 0 \\ \frac{3h_{Fx}}{m_{to}} \\ 0 \\ \frac{3h_{Fx}}{m_{to}} \end{bmatrix} V$$

System Identification



$$\begin{bmatrix} \dot{x}_{FA}(t) \\ \dot{x}_{FA}(t) \\ x_{SS}(t) \\ \dot{x}_{SS}(t) \end{bmatrix} = \begin{bmatrix} 0 & 1 & 0 & 0 \\ -\frac{k_{to}}{m_{to}} & \frac{81R}{32H^2} \frac{h_{Mz}}{m_{to}} - \frac{d_{to}}{m_{to}} - \frac{3h_{Fx}}{m_{to}} & 0 & 0 \\ 0 & 0 & 0 & 1 \\ 0 & -\frac{27R}{16H^2} \frac{h_{Fz}}{m_{to}} & -\frac{k_{to}}{m_{to}} & -\frac{d_{to}}{m_{to}} \end{bmatrix} \begin{bmatrix} x_{FA}(t) \\ \dot{x}_{FA}(t) \\ x_{SS}(t) \\ \dot{x}_{SS}(t) \end{bmatrix} + \begin{bmatrix} 0 \\ 0 \\ 0 \\ \frac{3}{2H} \frac{1}{m_{to}} \end{bmatrix} \begin{bmatrix} T_{gen} \\ \beta \end{bmatrix} + \begin{bmatrix} 0 \\ \frac{3h_{Fx}}{m_{to}} \\ 0 \\ \frac{3h_{Fz}}{m_{to}} \end{bmatrix} V$$

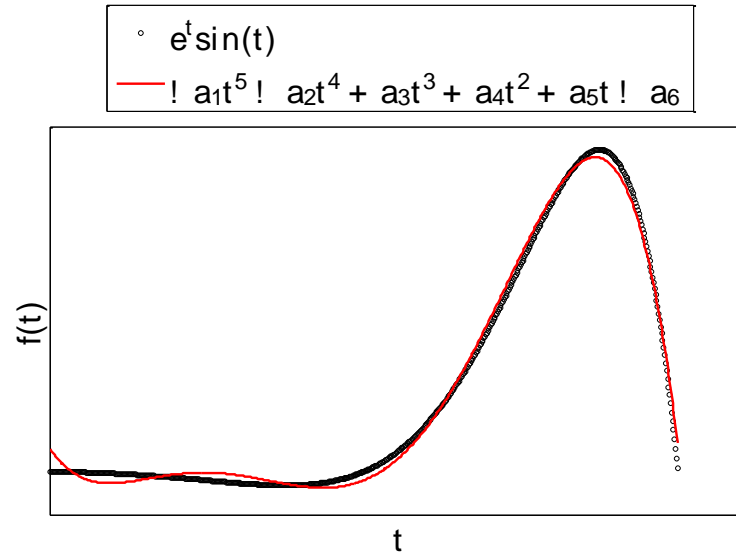
System Identification



Model $-a_1 t^5 - a_2 t^4 + a_3 t^3 + a_4 t^2 + a_5 t - a_6$

Structure / Form $-t^5 - t^4 + t^3 + t^2 + t - 1$ Parameters $a_1, a_2, a_3, a_4, a_5, a_6$

System Identification

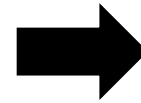
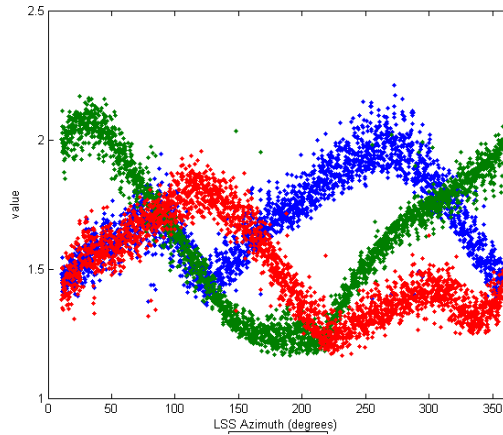
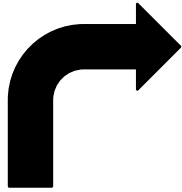


$$e^t * \frac{\sin}{\cos}$$

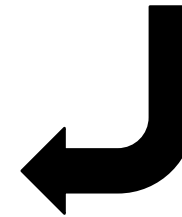
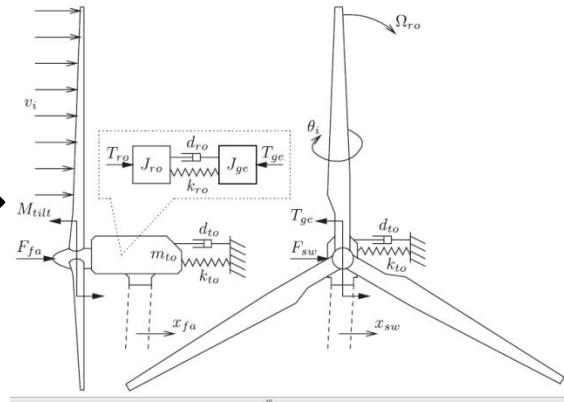
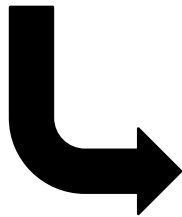
Symbolic Regression

$$e^t \sin(t)$$

Symbolic Regression



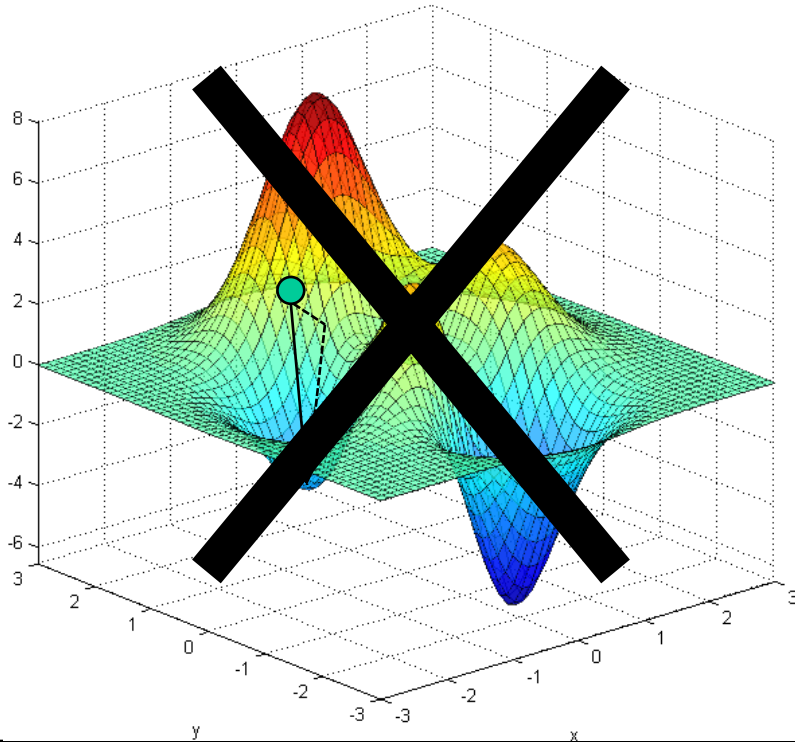
$$y = f(t, x, a, b, c)$$



Symbolic Regression

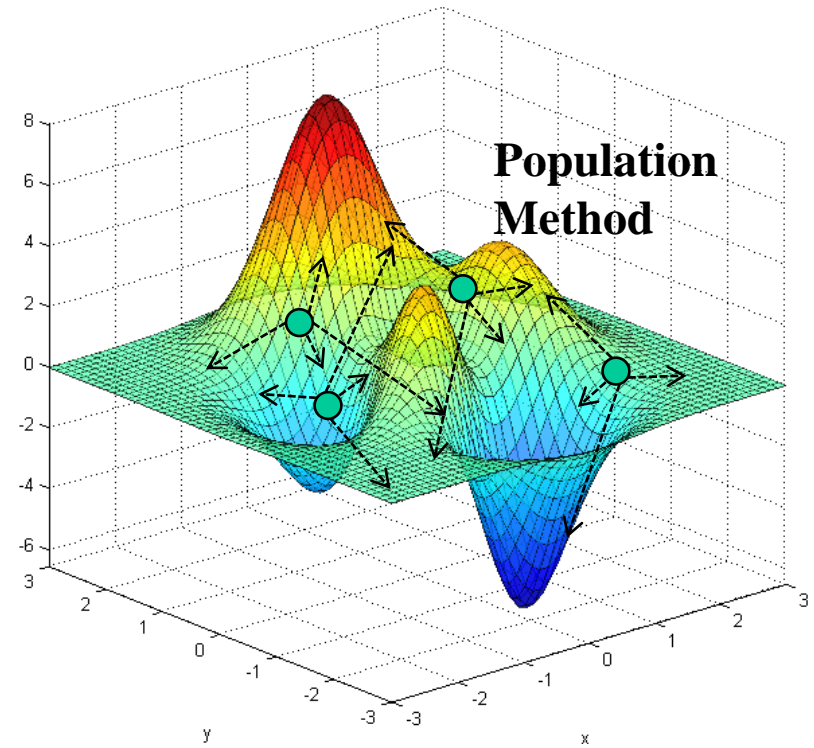
Gradient method

$$\vec{x} = \vec{x} + \alpha \nabla f(\vec{x})$$



Stochastic Method

$$\vec{x} = \textit{tweak}(\vec{x})$$



Genetic Programming

Initial Population

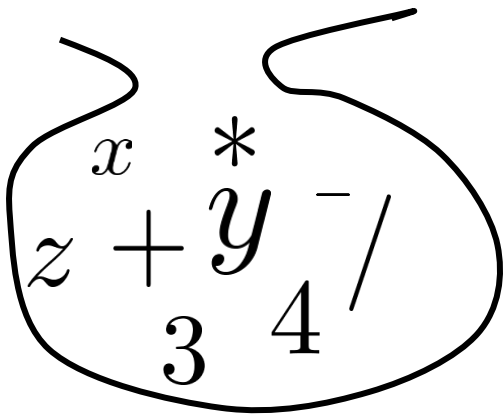
$$f_1 = x * y + z$$

$$f_2 = x/3 + y/z$$

$$f_3 = y - \frac{4}{(z+3)} + x$$

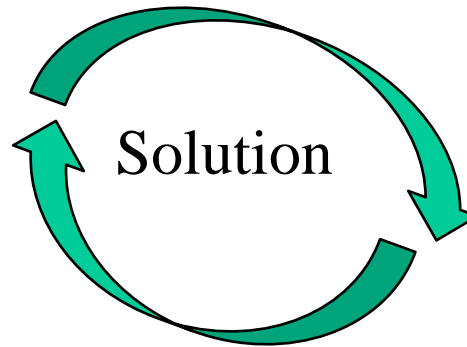
$$f_4 = y - z$$

$$f_5 = 3 * (x - y) + z^2$$



Fitness

$$E_i = \sum_{n=1}^N |f(n)^* - f_i(n)|$$



Selection

$$f_1 = x * y + z$$

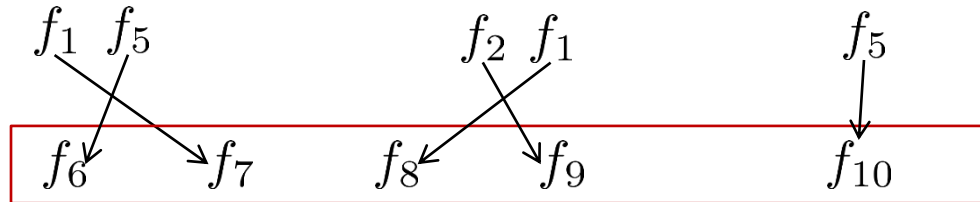
$$f_2 = x/3 + y/z$$

$$f_3 = y - \frac{4}{(z+3)} + x$$

$$f_4 = y - z$$

$$f_5 = 3 * (x - y) + z^2$$

Breeding



Genetic Programming - Representation

- Developmental
- Linear

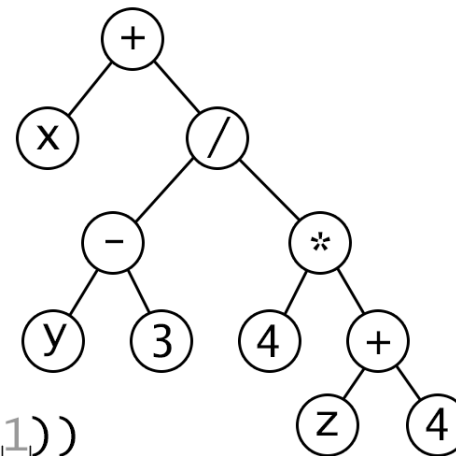
Genotype

```

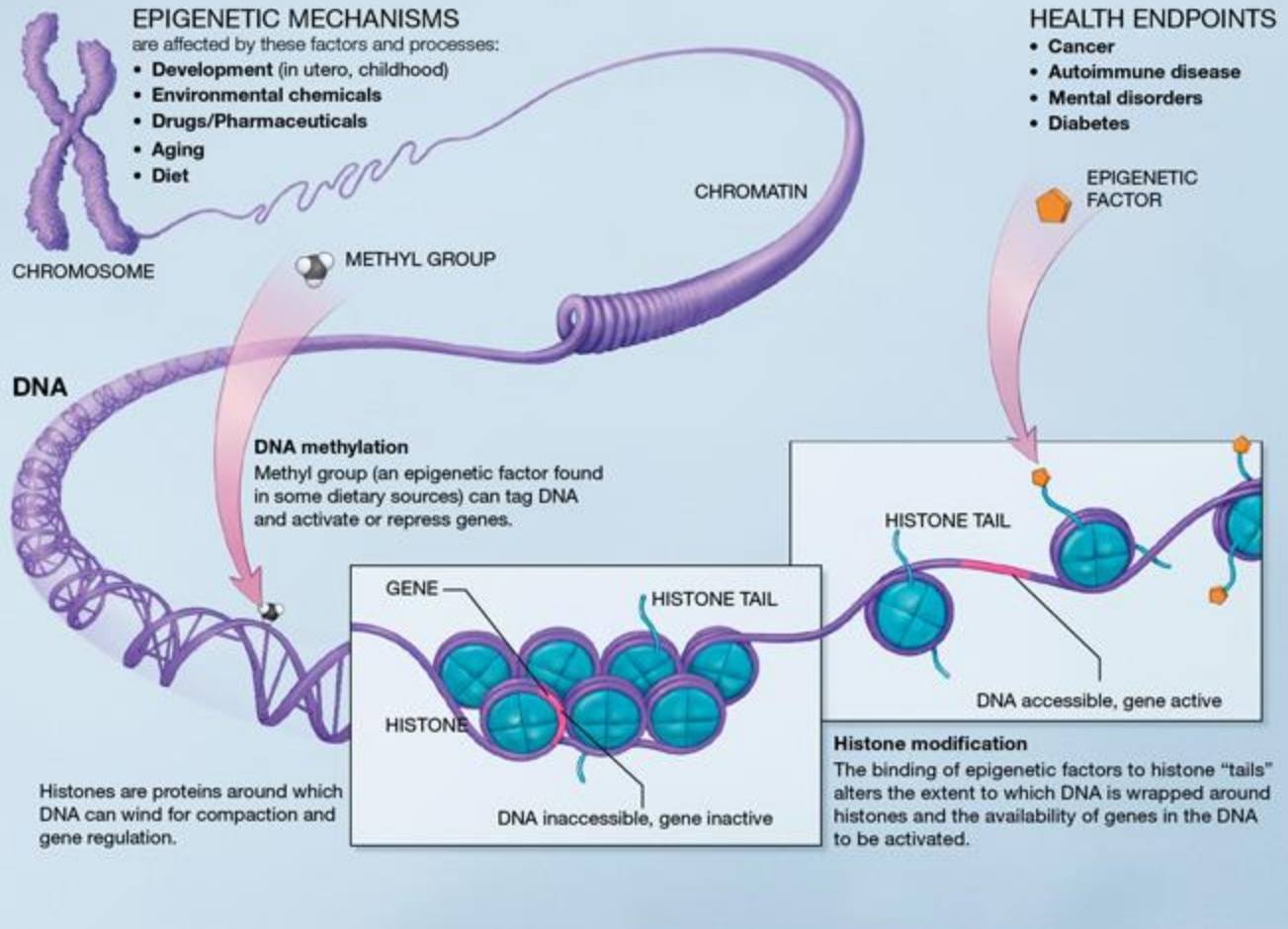
insert(x)
add
insert(y)
subR
insert(3)
UP
divR
insert(4)
mul
insert(z)
add
insert(4)
  
```

x
 $(x+0)$
 $(x+y)$
 $(x+(y-0))$
 $(x+(y-3))$
 $(x+(y-3))$
 $(x+(y-3)/1)$
 $(x+(y-3)/4)$
 $(x+(y-3)/(4*1))$
 $(x+(y-3)/(4z))$
 $(x+(y-3)/(4(z+0)))$
 $(x+(y-3)/(4(z+4)))$

Phenotype



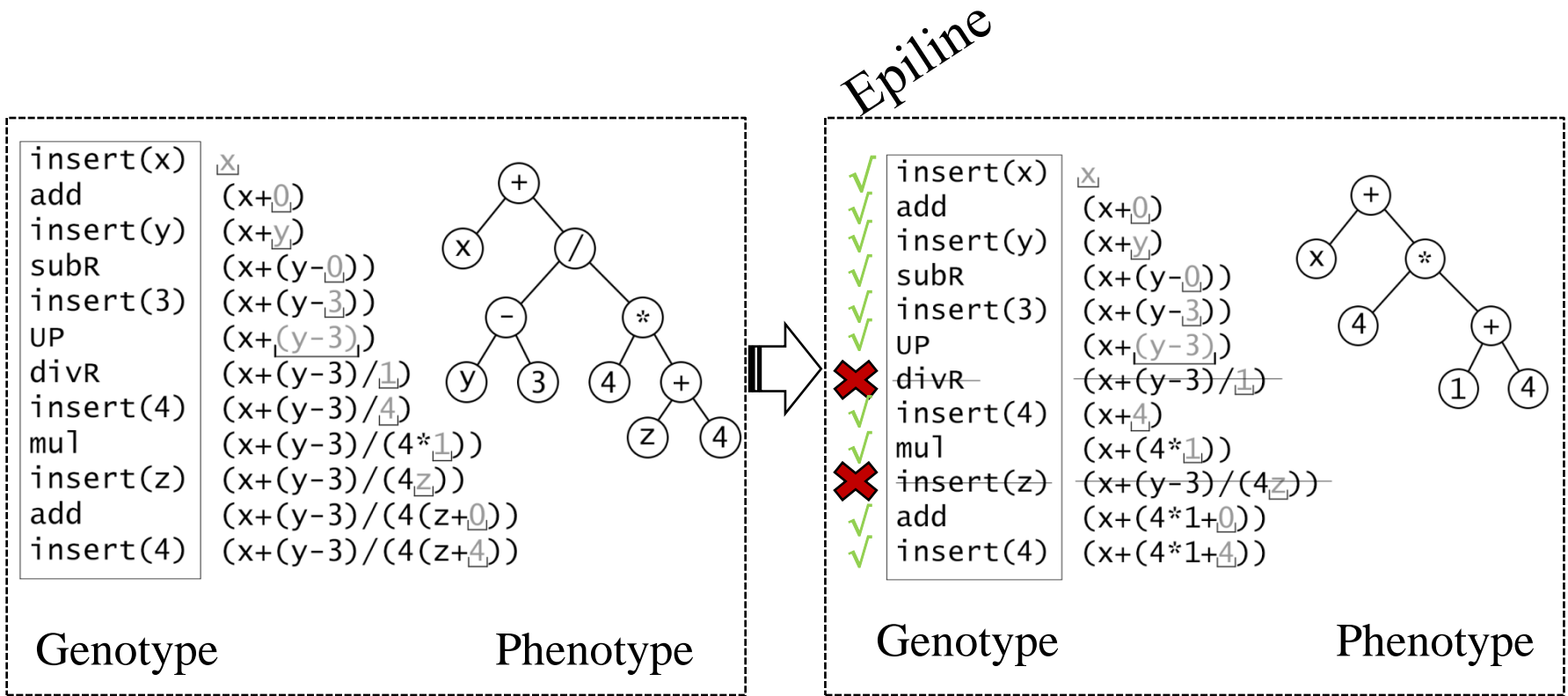
Epigenesis



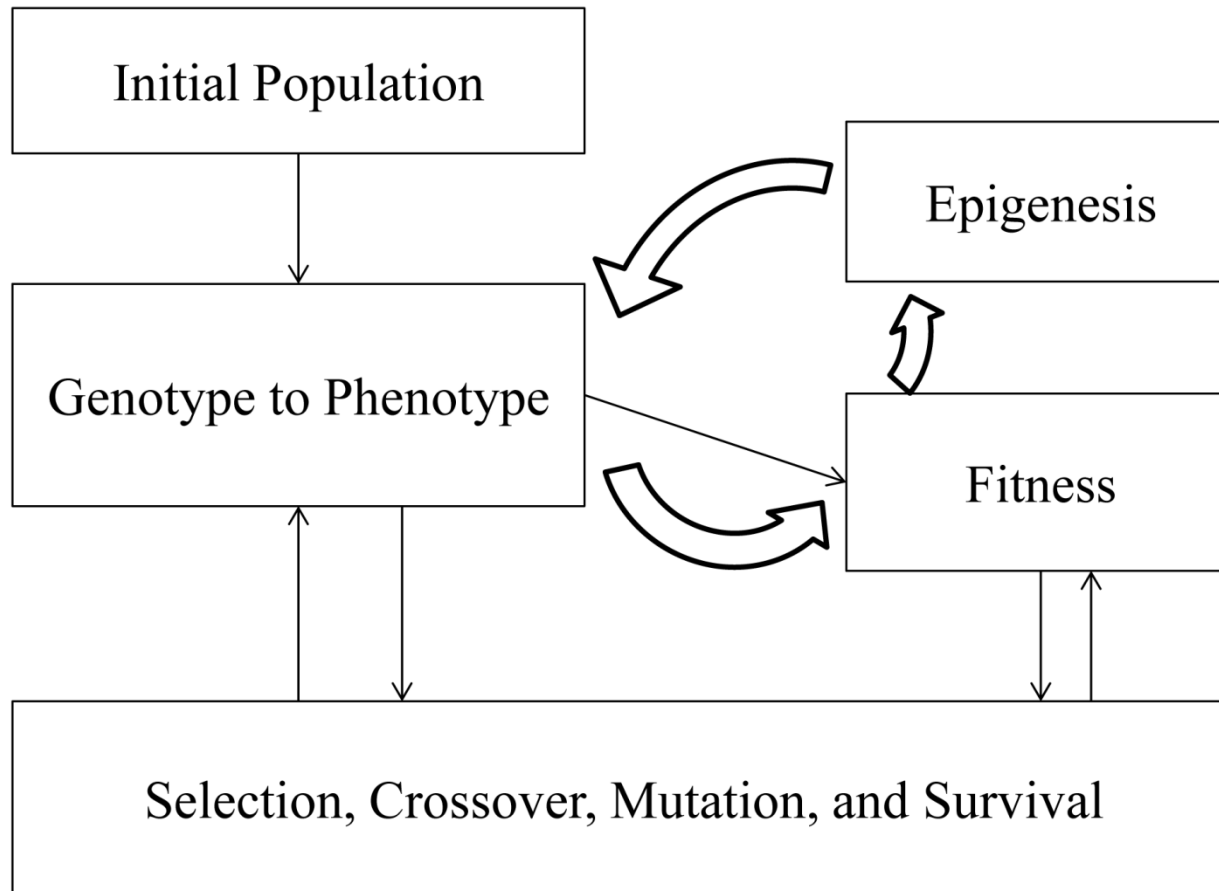
Environment

Inheritance

Genetic Programming - Epigenesis



Genetic Programming - Epigenesis



Genetic Programming - Epigenetic Hill Climber

```

gen ← genotype of individual
epi ← epigenetic line of individual
phen ← phenotype equation of individual
L ← length of genotype
sr ← switching rate
for number of hill climbing iterations do
  epiTemp ← epi
  for  $i \in L$  do
    if  $\text{rand}() < sr$  then
      /* flip the on/off state of marked
      indices in the epiline */
      epiTemp(i) ← !epi(i)
    end
  end
  /* get equation from genotype with updated
  gene expression */
  phenTemp ← GenToPhen(gen,epiTemp)
  /* update equation */
  if  $\text{fitness}(\text{phenTemp}) < \text{fitness}(\text{phen})$  then
    epi ← epiTemp
    phen ← phenTemp
    /* secondary size metric */
  else if  $\text{fitness}(\text{phenTemp}) == \text{fitness}(\text{phen})$  and
   $\text{size}(\text{phenTemp}) < \text{size}(\text{phen})$  then
    epi ← epiTemp
    phen ← phenTemp
  else
  end
end
end

```

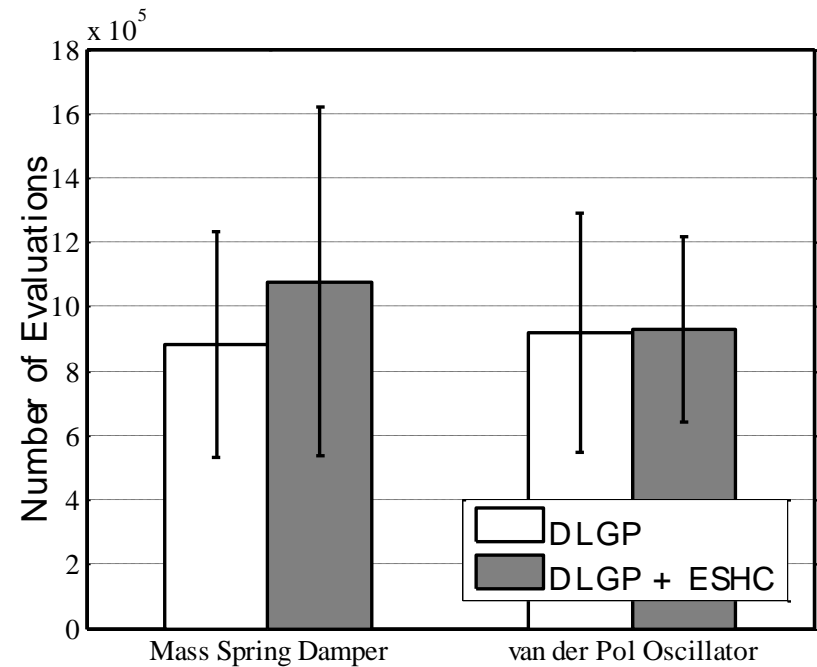
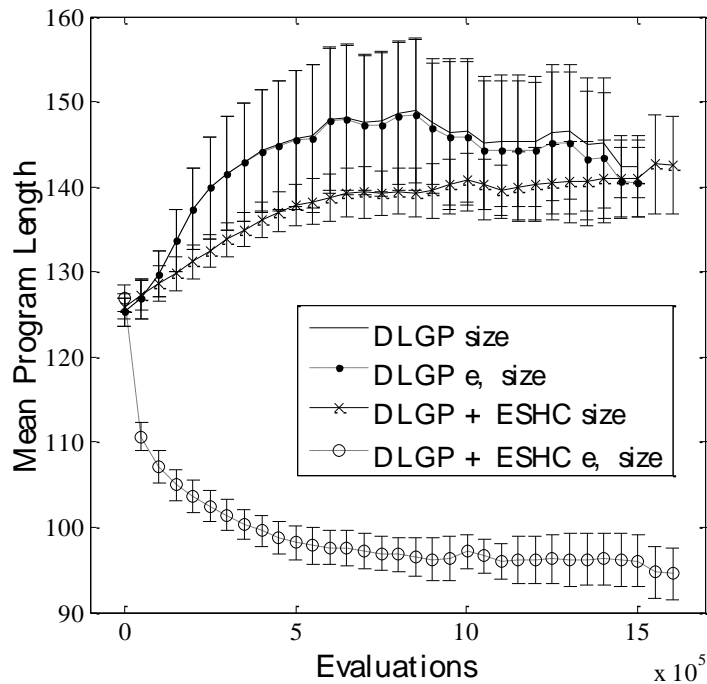
Algorithm 1: ESHC for one individual in population.

1. Randomly switch some genes on or off
2. Get new equation from genotype
3. If equation is better, update the epigenetic line

Examples

Table 1: Performance Comparisons

Method	Problem	Success Rate	Mean Evaluations	Mean Effective Program Size
DLGP	$\ddot{x} = -1/2(0.75\dot{x} + 3x - F)$	83.33%	8.83E+07	145.11
DLGP+ESHC		100%	10.77E+07	96.65
DLGP	$\ddot{x} = -1.5(x^2 + 1)\dot{x} - x$	83.33%	9.20E+05	140.06
DLGP+ESHC		100%	9.27E+05	101.34



Next Steps

NREL CART3

Table 1: Variable Definitions

State Variables	Other Variables	Control Inputs	Disturbances	Constants
x_{FA}	rotor azimuth, ϕ	T_{gen}	wind speed, V	tower mass, m_{to}
x_{SS}		blade pitch, β		rotor radius, R
\dot{x}_{FA}				hub height, H
\dot{x}_{SS}				
\ddot{x}_{FA}				
\ddot{x}_{SS}				

