

## Lab 2

### Gene Expression Algorithm

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LAB-2
import math, random
Gene Expression Algorithm
FUNCS = {'+':2, '-':2, '*':2, '/':2, 'sin':1, 'cos':1, 'exp':1,
        'log':1}
APPLY = {
    '+': lambda a,b: a+b, '-': lambda a,b: a-b,
    '*': lambda a,b: a*b, '/': lambda a,b: a if abs(b) < 1e-12
    else a/b, 'sin': lambda a: math.sin(a),
    'cos': lambda a: math.cos(a),
    'exp': lambda a: math.exp(max(-50, min(50, a))),
    'log': lambda a: math.log(abs(a+1))
}
TERMS = ['x', 'x']

def init_chrom(head, tail):
    g = random.choice(list(FUNCS) + TERMS) for _ in range(head)
    g = random.choice(TERMS) for _ in range(tail)
    c = random.uniform(-2, 2) for i, s in enumerate(g) if s == 'x'
    return f'g, g, "x": c'

def decode(g):
    seq = []
    need = 1
    i = 0
    while need > 0 and i < len(g):
        sym = g[i]
        seq.append(sym, i)
        need -= 1
        if sym in FUNCS:
            need += FUNCS[sym]
        i += 1
    return seq

def eval_expr(seq, x, c):
    spos = seq[0]
    if s in FUNCS:
        args = []
        for i in range(FUNCS[s]):

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        xj = eval_expr(seq[j], x, c); args.append(xj)
    try:
        return (FUNCS[s])(*args)
    except:
        return (float("inf"))
    return (x if s == "x" else c.get(seq, 0), seq)

def safe_eval(ch, x):
    try:
        v = eval_expr(ch, x, c)
    except:
        return float("inf")
    return v if math.isfinite(v) else float("inf")

def mse(ch, x, y):
    s = 0
    for xy in zip(x, y):
        d = safe_eval(ch, xy)
        if not math.isfinite(d):
            return float("inf")
        s += d * d
    return s / len(x)

def mutate(ch, head, tail, pm = 0.05, pc = 0.1):
    g = ch["g"]
    c = ch["c"]
    for i in range(n):
        if random.random() < pm:
            g[i] = random.choice(list(FUNCS) + TERMS)
            if i < head else TERMS
            if g[i] == "x":
                c[i] = random.uniform(-2, 2)
            elif i < len(c):
                c[i] = c[i]
        for i in list(c):
            if random.random() < pc:
                c[i] = c[i] + random.gauss(0, 0.1)

def crossover(p1, p2):
    n = len(p1["g"])
    cut = random.randint(1, n-1)
    def make(g1, p1):
        g = p1["g"][:cut] + p2["g"][cut:]
        for i, s in enumerate(g):
            if s == "x":
                c[i] = p1["c"] if i < cut else p2["c"]
            else:
                c[i] = random.uniform(-2, 2)

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    return {"g": g, "c": c}
    return make(a,b), make(b,a)

def select(pop, fit, k=3):
    i = min(random.sample(range(len(pop)), k), key=lambda j: fit[j])
    return {"g": pop[i]["g"][:], "c": dict(pop[i]["c"])}

def evolve(f, target=(-3,3), pts=64, popn=100, head=10, gens=100):
    cx = 0.7; seed = None:
    if seed: random.seed(seed)
    tail = head * (max(functools.values()) - 1) + 1
    X = [target[0] + (target[1] - target[0]) * i / (pts - 1) for i in range(pts)]
    Y = [f(x) for x in X]
    pop = [init_chrom(head, tail) for _ in range(popn)]
    best, fit = float("inf"), None
    for gen in range(gens):
        fits = [mse(ind, X, Y) for ind in pop]
        b = min(range(popn), key=lambda i: fits[i])
        if fits[b] < best: best, fit = fits[b], {"g": pop[b]["g"][:], "c": dict(pop[b]["c"])}
        new = [fit]
        while len(new) < popn:
            p1 = select(pop, fits); p2 = select(pop, fits)
            c1, c2 = crossover(p1, p2) if random.random() < cx else (p1, p2)
            mutate(c1, head, tail); mutate(c2, head, tail)
            new += [c1, c2]
        pop = new
        if (gen+1) % 20 == 0: print("Gen", gen+1, "Best", best)
    return fit, best

if __name__ == "__main__":
    f = lambda x: x**3 - 0.5 * x + math.sin(x)
    best, err = evolve(f, gens=100)
    print("Best error:", err)
    print("Seeds:", [safe_eval(best, x) for x in [-2, -1, 0, 1, 2]])

```

Step 1: Fitness function:  $f(x) = x^2$   
 Encoding technique: 0 to 3  
 Use chromosome of fixed length (genotype)

Step 2: Initial population

S. No.	(Genotype) Initial chromosome	Phenotype (express <sup>n</sup> )	Value	Fitness	P
1	+xx	$x^2$	12	144	0.1247
2	+xx	$2x$	25	625	0.5411
3	x	x	5	25	0.0216
4	-x2	$x^2$	19	361	0.3125
Sum				1155	
avg				288.75	
max				625	

Actual count	Expected count
1	0.5
2	2.1
0	0.08
1	1.25

Step 3: Selection of mating pool

S. No.	Selected chromosome	Crossover point	Offspring	Phenotype
1	+xx	2	*x+	$x+(x+...)$
2	+xx	1	+xx	$2x$
3	+xx	3	+x-	$x+(x-...)$
4	-x2	1	+x2	$x+2$

  

x value	Fitness
13	169
24	576
27	729
17	289

Step 4: Crossover = perform crossover randomly chosen gene position  
 (not saw bits)  
 max fitness after crossover = 729

Step 5: Mutation

S. No.	Offspring before mutation	Mutation applied	Offspring after mutation	Phenotype
1	+x+	+ → -	*x-	$x+(x-...)$
2	+x2	none	+x2	$2x$
3	+x-	- → +	-x+	$x+(x+...)$
4	+x2	none	+x2	$x+2$

  

x value	Fitness
29	841
24	576
27	729
20	400

Step 6: Gene Expression and evaluation

decode each genotype → phenotype

calculate fitness

$$\sum f(x) = 841 + 576 + 729 + 400 = 2546$$

$$\text{avg} = 636.5$$

$$\text{max} = 841$$

Step 7: Iterate until convergence

Repeat step 3 to 6 until fitness improvement is negligible  
 or generation limit has reached

Output:

1000 generations

Genes: [29.53, 29.82, 29.84, 28.57, 16.09, 21.83, 23.83, 30.31, 28.51, 26.22]

$$x = 26.37$$

$$f(x) = 695.45$$

Code:

```
print("Shreya Raj 1BM23CS317")

import math, random

# --- Function set ---
FUNCS = {"+":2, "-":2, "*":2, "/":2, "sin":1, "cos":1, "exp":1, "log":1}
APPLY = {
    "+":lambda a,b:a+b, "-":lambda a,b:a-b, "*":lambda a,b:a*b,
    "/":lambda a,b:a if abs(b)<1e-12 else a/b,
    "sin":lambda a:math.sin(a), "cos":lambda a:math.cos(a),
    "exp":lambda a:math.exp(max(-50,min(50,a))),
    "log":lambda a:math.log(abs(a)+1)
}
TERMS=["x", "C"]

def init_chrom(head,tail):
    g=[random.choice(list(FUNCS)+TERMS) for _ in range(head)]
    g+=[random.choice(TERMS) for _ in range(tail)]
    c={i:random.uniform(-2,2) for i,s in enumerate(g) if s=="C"}
    return {"g":g, "c":c}

def decode(g):
    seq=[];need=1;i=0
    while need>0 and i<len(g):
        sym=g[i];seq.append((sym,i));need-=1
        if sym in FUNCS: need+=FUNCS[sym]
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i+=1  
return seq
```

```
def eval_expr(seq,i,x,c):  
    s,pos=seq[i]  
    if s in FUNCS:  
        args=[];j=i+1  
        for _ in range(FUNCS[s]):  
            v,j=eval_expr(seq,j,x,c);args.append(v)  
        try:return (APPLY[s](*args),j)  
        except: return (float("inf"),j)  
    return (x if s=="x" else c.get(pos,0),i+1)
```

```
def safe_eval(ch,x):  
    try:v,_=eval_expr(decode(ch["g"]),0,x,ch["c"])  
    except:return float("inf")  
    return v if math.isfinite(v) else float("inf")
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def mse(ch,X,Y):  
    s=0  
    for x,y in zip(X,Y):  
        d=safe_eval(ch,x)-y  
        if not math.isfinite(d): return 1e12  
        s+=d*d  
    return s/len(X)
```

```

def mutate(ch,head,tail,pm=0.05,pc=0.1):
    g,c=ch["g"],ch["c"];n=len(g)
    for i in range(n):
        if random.random()<pm:
            g[i]=random.choice((list(FUNCS)+TERMS) if i<head else TERMS)
            if g[i]=="C":c[i]=random.uniform(-2,2)
            elif i in c:del c[i]
    for i in list(c):
        if random.random()<pc:c[i]+=random.gauss(0,0.1)

def crossover(a,b):
    n=len(a["g"]);cut=random.randint(1,n-1)
    def make(pa,pb):
        g=pa["g"][:cut]+pb["g"][cut:];c={}
        for i,s in enumerate(g):
            if s=="C":c[i]=(pa["c"] if i<cut else pb["c"]).get(i,random.uniform(-2,2))
        return {"g":g,"c":c}
    return make(a,b),make(b,a)

def select(pop,fit,k=3):
    i=min(random.sample(range(len(pop)),k),key=lambda j:fit[j])
    return {"g":pop[i]["g"][:], "c":dict(pop[i]["c"])}

def evolve(f,target=(-3,3),pts=64,popn=100,head=10,gens=100,cx=0.7,seed=None):
    if seed:random.seed(seed)
    tail=head*(max(FUNCS.values())-1)+1

```



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X=[target[0]+(target[1]-target[0])*i/(pts-1) for i in range(pts)]
Y=[f(x) for x in X]
pop=[init_chrom(head,tail) for _ in range(popn)]
best,fit=float("inf"),None
for gen in range(gens):
    fits=[mse(ind,X,Y) for ind in pop]
    b=min(range(popn),key=lambda i:fits[i])
    if fits[b]<best:best,fit=fits[b],{"g":pop[b]["g"][:],"c":dict(pop[b]["c"])}
    new=[fit]
    while len(new)<popn:
        p1=select(pop,fits);p2=select(pop,fits)
        c1,c2=crossover(p1,p2) if random.random()<cx else (p1,p2)
        mutate(c1,head,tail);mutate(c2,head,tail)
        new+=[c1,c2]
    pop=new
    if (gen+1)%20==0:print("Gen",gen+1,"Best",best)
return fit,best

```

```

if __name__=="__main__":
    f=lambda x:x**3-0.5*x+math.sin(x)
    best,err=evolve(f,gens=100)
    print("Best error:",err)
    print("Preds:",[safe_eval(best,x) for x in [-2,-1,0,1,2]])

```

Output:

```
➦ Shreya Raj 1BM23CS317  
Gen 20 Best 0.2711388144567344  
Gen 40 Best 0.08276309290483551  
Gen 60 Best 0.02638736211355352  
Gen 80 Best 0.0038234107968136122  
Gen 100 Best 0.0035796711491496228  
Best error: 0.0035796711491496228  
Preds: [-7.947649551427884, -1.2578413596603832, 0.0, 1.2578413596603832, 7.947649551427884]
```