

Note: Updated Maple takes 2D form  $\frac{a}{b}$  rather than a/b

## Final

1.  $x(n) = 2x(n-1) - x(n-3)$ ,  $x(0)=1$ ,  $x(1)=1$ ,  $x(2)=2$

Use Orbk to find long term behavior of 3<sup>rd</sup> order system

evalf( $\text{Orbk}(3, z, 2 \cdot z[1] - z[3], [1, 1, 2], 1000, 1001)[2]$ )  $\leftarrow$  Orbk sets the value of  $x(0)$  at k=1  
 $\text{Orbk}(3, z, 2 \cdot z[1] - z[3], [1, 1, 2], 1000, 1001)[1]$   
= 1.618033989

2.  $x'(t) = \frac{5}{2}x(t)(1-x(t))(1-\frac{1}{2}x(t))$

a) Find underlying function

$$f(x) = \frac{5}{2}x(1-x)(1-\frac{1}{2}x)$$

By hand: solve where rate of change is 0

$$0 = \frac{5}{2}x(1-x)(1-\frac{1}{2}x)$$

$$x=0, x=1, x=2$$

In Maple: EQuP([ $\frac{5}{2} \cdot x \cdot (1-x) \cdot (1-\frac{1}{2}x)$ ], [x])  $\Rightarrow \{[0], [1], [2]\}$

b) Find derivative of underlying function

$$f'(x) = (\frac{5}{2}x - \frac{5}{2}x^2)(1-\frac{1}{2}x)$$

$$= \frac{5}{2}x - \frac{5}{4}x^2 - \frac{5}{2}x^2 + \frac{5}{4}x^3$$

$$f'(x) = \frac{5}{2}x - \frac{15}{4}x^2 + \frac{5}{4}x^3$$

$$f'(x) = \frac{5}{2} - \frac{15}{2}x + \frac{15}{4}x^2$$

Check derivative values at equilibrium points

$$f'(0) = \frac{5}{2} \Rightarrow \text{unstable } \frac{5}{2} > 0$$

$$f'(1) = -\frac{5}{4} \Rightarrow \text{stable } -\frac{5}{4} < 0 \Rightarrow x=1 \text{ is the stable equilibrium point}$$

$$f'(2) = \frac{5}{2} \Rightarrow \text{unstable } \frac{5}{2} > 0$$

In Maple: SEquP([ $\frac{5}{2} \cdot x(1-x)(1-\frac{1}{2}x)$ ], [x])  $\Rightarrow \{[1]\}$

c)  $x(100) = 1$  because 1 is a stable equilibrium therefore at initial values around 1, the system will converge to 1

Checked in Maple:

$$\text{evalf}(\text{subs}(t=100, \text{dsolve}(\{\text{diff}(x(t), t) = \frac{5}{2} \cdot x(t) \cdot (1-x(t)) \cdot (1-\frac{1}{2} \cdot x(t)), x(0)=0.1\}, x(t))))$$
$$\Rightarrow x(100) = 0.9999999999 \Rightarrow \text{approximately } 1$$

$$3. \alpha(n) = \frac{5}{2}\alpha(n-1)(1-\alpha(n-1))(1-\frac{1}{2}\alpha(n-1))$$

a) Find fixed points of underlying function (where  $\alpha = F(\alpha)$ )

$$F = \frac{5}{2}\alpha(1-\alpha)(1-\frac{1}{2}\alpha)$$

$$\alpha = \frac{5}{2}\alpha(1-\alpha)(1-\frac{1}{2}\alpha)$$

$$\text{Maple: evalf}(FP([\frac{5}{2} \cdot \alpha \cdot (1-\alpha) \cdot (1-\frac{1}{2} \cdot \alpha)], [\alpha])) \Rightarrow \{[0], [0.475304923], [2.524695077]\}$$

$$\alpha = 0, \alpha = 0.475304923, \alpha = 2.524695077$$

b) Find which equilibrium solution satisfies  $|F'(\alpha)| < 1$

$$F' = \frac{5}{2} - \frac{15}{2}\alpha + \frac{15}{4}\alpha^2$$

$$F'(0) = \frac{5}{2} \Rightarrow \text{unstable } |\frac{5}{2}| > 1$$

$$F'(0.475304923) = -0.217606535645 \Rightarrow \text{stable } |-0.217606535645| < 1$$

$$F'(2.524695077) = 7.46760654186 \Rightarrow \text{unstable } |7.46760654186| > 1$$

$$\text{In Maple: evalf}(SFP([\frac{5}{2} \cdot \alpha \cdot (1-\alpha) \cdot (1-\frac{1}{2} \cdot \alpha)], [\alpha])) \Rightarrow [0.4753049232]$$

$$\alpha = 0.4753049232 \text{ is the stable equilibrium}$$

c)  $\alpha(1000) = 0.4753049232$  because the system will converge to the stable equilibrium if it begins in the neighborhood of that equilibrium.

$$\text{Checked in Maple: Orb}([\frac{5}{2} \cdot \alpha \cdot (1-\alpha) \cdot (1-\frac{1}{2} \cdot \alpha)], [\alpha], [0.1], 1000, 1000)[1] \Rightarrow [0.4753049232]$$

4. If there are equal proportions of AA, Aa, and aa, then  $u=v=w=\frac{1}{3}$  because

$$u+v+w=1.$$

✓ Generation 1 is  $k=0$

$$a) \text{Orb}(HW3(u, v, w), [u, v, w], [\frac{1}{3}, \frac{1}{3}, \frac{1}{3}], 1, 1)[1][2]$$

$$\Rightarrow \frac{1}{2} \Rightarrow \text{genotype Aa proportion} = \frac{1}{2}$$

$$b) \text{Orb}(HW3(u, v, w), [u, v, w], [\frac{1}{3}, \frac{1}{3}, \frac{1}{3}], 999, 999)[1][2]$$

$$\Rightarrow \frac{1}{2} \Rightarrow \text{genotype Aa proportion} = \frac{1}{2}$$

5. If the probabilities of mating are unequal then you must incorporate the probability matrix and multiply each mating with the probability of that match occurring

a)  $\text{Orb}(\text{HW3g}(u, v, w, [[.1, .2, .1], [.1, .1, .1], [.1, .1, .1]]), [u, v, w], [\frac{1}{3}, \frac{1}{3}, \frac{1}{3}], 1, 1)[1][2]$   
 $\Rightarrow 0.5 \Rightarrow \text{proportion of genotype Aa} = 0.5$

Used any probability matrix where AA (female)  $\times$  Aa (male) was double that of the other crosses

b)  $\text{Orb}(\text{HW3g}(u, v, w, [[.1, .2, .1], [.1, .1, .1], [.1, .1, .1]]), [u, v, w], [\frac{1}{3}, \frac{1}{3}, \frac{1}{3}], 999, 999)[1][2]$   
 $\Rightarrow 0.3974661806 \Rightarrow \text{proportion of genotype Aa} = 0.3974661806$

6. Use underlying transformation to examine long term behavior

$$(x, y) \rightarrow \left( \frac{1+x+4y}{2+x+3y}, \frac{1+2x+3y}{3+x+2y} \right)$$

$$\text{OrbF}\left([\frac{1+x+y}{2+x+3y}, \frac{1+2x+3y}{3+x+2y}], [x, y], [100, 1000], 1000, 1001\right)[1][2] \Rightarrow 0.7478789080$$

$$\text{OrbF}\left([\frac{1+x+y}{2+x+3y}, \frac{1+2x+3y}{3+x+2y}], [x, y], [100, 1000], 1000, 1001\right)[2][2] \Rightarrow 0.7478789080$$

Equilibrium of y at 0.7478789080

7. Find how many susceptible and infected in the long run and then do  $N - s - i$  to find recovered ( $1000 - s - i$ )

a)  $\text{SEquP}(\text{SIRS}(s, i, 0.05, 0.5, 100, 1000), [s, i]) \Rightarrow \{[1000, 0]\}$

OR  $\text{Dis}(\text{SIRS}(s, i, 0.05, 0.5, 100, 1000), [s, i], [300, 300], 0.01, 20)[-1]$   
 $\Rightarrow [20.01, [999.9666301, 6.997984915 \times 10^{-686}]] \Rightarrow \text{approx. } [1000, 0]$

$$1000 - 1000 - 0 = 0 \text{ recovered}$$

b)  $\text{SEquP}(\text{SIRS}(s, i, 14, 0.5, 100, 1000), [s, i]) \Rightarrow \{[71.42857143, 4.619758351]\}$

$$1000 - 71.42857143 - 4.619758351 = 923.951670219 \text{ recovered}$$

c) Find value of b where  $i=0 \Rightarrow b = 0.1$

in  $\begin{cases} \text{SIRS}(s, i, b, 0.5, 100, 1000) \\ \text{SysEq} := \{-sib + 500.0 - 0.5s - 0.5i = 0, sib - 100i = 0\} \end{cases}$  from SIRS model  
 solve for b

8. a) SEquP(GeneNet(0, 1, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3])

Stable equilibrium at 0.6823278038

b) Time Series(GeneNet(0, 3, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.2, 0.1, 0.3, 0.1, 0.4, 0.5], 0.01, 200, 1)

⇒ graph in Maple ⇒ there is a horizontal asymptote

SEquP(GeneNet(0, 3, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3])

Height of horizontal asymptote = stable equilibrium at 1.213411663

c) Found where the system no longer had a stable equilibrium

First found where SEquP outputted an empty set with step size 1 between 3 and 50. This gave a = 8 as the least entry with an empty set.

Then found where the output was the empty set between 7 and 8 with step size 0.1: 7.4. Then used step size 0.01 and the greatest a with a nonempty set was 7.39. Code in Maple.

9. Found stable equilibrium point of Chemostat model

SEquP(ChemoStat(N, C, 2.5, 2.7), [N, C]) ⇒ { [5.083333333, 0.6666666667] }

a) Bacterial population density = 5.083333333

Bacterial pop density

b) Nutrient concentration = 0.6666666667

Nutrient concentration

10. Created the transition matrix for the probability of moving from page to page

Rows of transition matrix must sum to 1

Called it matrix P in Maple

Take the limit:  $\lim_{n \rightarrow \infty} P^n$  or in Maple  $P^{1000}$

Full code in Maple

Site 1: 0.07692307655 Site 9: 0.1538461530

```
> #Shreya Ghosh, Maple Code for Final Exam
> read "/Users/shreyaghosh/Documents/DMB.txt"
      First Written: Nov. 2021
```

*This is DMB.txt, A Maple package to explore Dynamical models in Biology (both discrete and continuous)*

*accompanying the class Dynamical Models in Biology, Rutgers University. Taught by Dr. Z. (Doron Zeilberger)*

*The most current version is available on WWW at:*

*<http://sites.math.rutgers.edu/~zeilberg/tokhniot/DMB.txt>.*

*Please report all bugs to: DoronZeil at gmail dot com .*

*For general help, and a list of the MAIN functions,  
type "Help();". For specific help type "Help(procedure\_name);"*

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*For a list of the supporting functions type: Help1();*

*For help with any of them type: Help(ProcedureName);*

---

*For a list of the functions that give examples of Discrete-time dynamical systems (some famous),  
type: HelpDDM();*

*For help with any of them type: Help(ProcedureName);*

---

*For a list of the functions continuous-time dynamical systems (some famous) type: HelpCDM();*

*For help with any of them type: Help(ProcedureName);*

(1)

```
> #1)
> evalf(Orbk(3,z,2·z[1] - z[3], [1, 1, 2], 1000, 1001)[2])
      1.618033989
```

(2)

```
> #2a)
> EquP([5/2 · x · (1 - x) · (1 - 1/2 · x)], [x])
      {[0], [1], [2]}
```

(3)

```

> #2b)
> SEquP $\left(\left[\frac{5}{2} \cdot x \cdot (1 - x) \cdot \left(1 - \frac{1}{2} \cdot x\right)\right], [x]\right)$ 

$$\{[1.\}\}$$
 (4)

> #2c)
> evalf $\left(\text{subs}\left(t = 100, \text{dsolve}\left(\left\{\text{diff}(x(t), t) = \frac{5}{2} \cdot x(t) \cdot (1 - x(t)) \cdot \left(1 - \frac{1}{2} \cdot x(t)\right), x(0) = 0.1\right\}, x(t)\right)\right)\right)$ 

$$x(100) = 0.9999999999$$
 (5)

>
> #3a)
> evalf $\left(\text{FP}\left(\left[\frac{5}{2} \cdot x \cdot (1 - x) \cdot \left(1 - \frac{1}{2} \cdot x\right)\right], [x]\right)\right)$ 

$$\{[0.], [0.475304923], [2.524695077]\}$$
 (6)

> #3b)
> evalf $\left(\text{SFP}\left(\left[\frac{5}{2} \cdot x \cdot (1 - x) \cdot \left(1 - \frac{1}{2} \cdot x\right)\right], [x]\right)\right)$ 

$$\{[0.475304923]\}$$
 (7)

> #3c)
> Orb $\left(\left[\frac{5}{2} \cdot x \cdot (1 - x) \cdot \left(1 - \frac{1}{2} \cdot x\right)\right], [x], [0.1], 1000, 1000\right)[1]$ 

$$[0.4753049232]$$
 (8)

>
> #4a)
> Orb $\left(HW3(u, v, w), [u, v, w], \left[\frac{1}{3}, \frac{1}{3}, \frac{1}{3}\right], 1, 1\right)[1][2]$ 

$$\frac{1}{2}$$
 (9)

> #4b)
> Orb $\left(HW3(u, v, w), [u, v, w], \left[\frac{1}{3}, \frac{1}{3}, \frac{1}{3}\right], 999, 999\right)[1][2]$ 

$$\frac{1}{2}$$
 (10)

>
> #5a)
> Orb $\left(HW3g(u, v, w, [[.1, .2, .1], [.1, .1, .1], [.1, .1, .1]]), [u, v, w], \left[\frac{1}{3}, \frac{1}{3}, \frac{1}{3}\right], 1, 1\right)[1][2]$ 

$$0.5000000001$$
 (11)

> Orb $\left(HW3g(u, v, w, [[.1, .2, .1], [.1, .1, .1], [.1, .1, .1]]), [u, v, w], \left[\frac{1}{3}, \frac{1}{3}, \frac{1}{3}\right], 999,$ 

```

=> #6)  
=>  $OrbF\left(\left[\frac{1+x+y}{2+x+3 \cdot y}, \frac{1+x+3 \cdot y}{3+x+2 \cdot y}\right], [x, y], [100, 1000], 1000, 1001\right)[1][2]$   
0.7478789080 (13)

$$\text{OrbF}\left(\left[\frac{1+x+y}{2+x+3 \cdot y}, \frac{1+x+3 \cdot y}{3+x+2 \cdot y}\right], [x, y], [100, 1000], 1000, 1001\right)[2][2] \\ \textcolor{blue}{0.7478789080} \quad (14)$$

```
> #7a)
> SEquP(SIRS(s, i, 0.05, 0.5, 100, 1000), [s, i])
      {[1000., 0.]} (15)
```

$$\text{Dis}(SIRS(s, i, 0.05, 0.5, 100, 1000), [s, i], [300, 300], 0.01, 20)[-1] \\ [20.01, [999.9666301, 6.997984915 \times 10^{-686}]] \quad (16)$$

> #7b)  
 >  $SEquP(SIRS(s, i, 1.4, 0.5, 100, 1000), [s, i])$   
 $\quad \{[71.42857143, 4.619758351]\}$  (17)

=> #7c)  
=>  $SIRS(s, i, b, 0.5, 100, 1000)$   
= [  $-b s i + 500.0 - 0.5 s - 0.5 i, b s i - 100 i$  ] (18)

```
> SysEq := { -b s i + 500.0 - 0.5 s - 0.5 i = 0, b s i - 100 i = 0 }
           SysEq := { -b s i + 500.0 - 0.5 s - 0.5 i = 0, b s i - 100 i = 0 } (19)
=>
> solve(SysEq, {s, i});
```

$$\{i=0., s=1000.\}, \left\{i = \frac{0.4975124378 (10.b - 1.)}{b}, s = \frac{100.}{b}\right\} \quad (20)$$

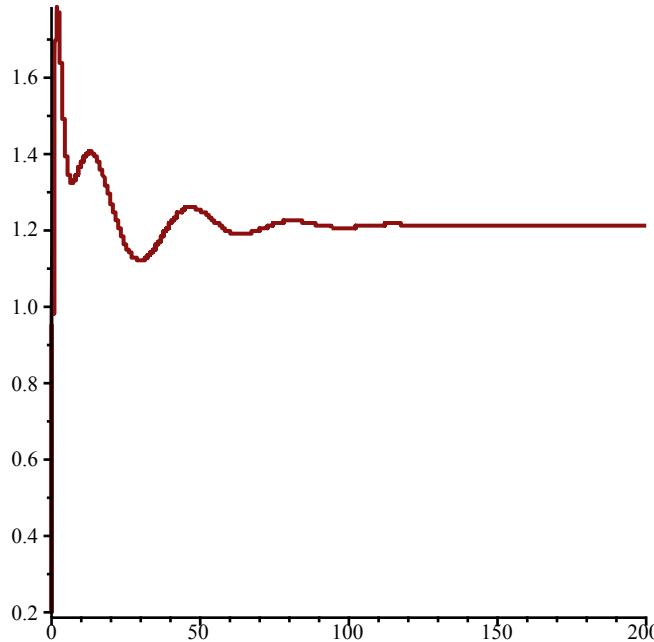
$$> \text{solve}\left(0 = \frac{0.4975124378(10.b - 1.)}{b}, b\right) \\ = 0.1000000000 \quad (21)$$

>  $\text{solve}\left(1000 = \frac{100}{b}, b\right)$  0.1000000000 (22)

```
> #8a)
> SEquP(GeneNet(0, 1, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3])
{[0.6823278038, 0.6823278038, 0.6823278038, 0.6823278038, 0.6823278038, 0.6823278038]} (23)
```

```
> #8b)  
>  
> TimeSeries(GeneNet(0, 3, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3], [0.2, 0.1,
```

```
0.3, 0.1, 0.4, 0.5 ], 0.01, 200, 1)
```



```
> SEquP(GeneNet(0, 3, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3])  
 {[1.213411663, 1.213411663, 1.213411663, 1.213411663, 1.213411663, 1.213411663]} (24)
```

```
> #8c)
```

```
> for a from 3 by 1 to 50
```

```
do
```

```
print(a);
```

```
SEquP(GeneNet(0, a, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3]);
```

```
end do;
```

3

```
{[1.213411663, 1.213411663, 1.213411663, 1.213411663, 1.213411663, 1.213411663]}
```

4

```
{[1.378796700, 1.378796700, 1.378796700, 1.378796700, 1.378796700, 1.378796700]}
```

5

```
{[1.515980228, 1.515980228, 1.515980228, 1.515980228, 1.515980228, 1.515980228]}
```

6

```
{[1.634365293, 1.634365293, 1.634365293, 1.634365293, 1.634365293, 1.634365293]}
```

7

```
{[1.739203861, 1.739203861, 1.739203861, 1.739203861, 1.739203861, 1.739203861]}
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∅ (25)

> **for** *a* **from** 7 **by** 0.1 **to** 8  
do  
*print(a);*

```

 $SEquP(GeneNet(0, a, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3]) ;$ 
end do;
    7
{[1.739203861, 1.739203861, 1.739203861, 1.739203861, 1.739203861, 1.739203861]}

    7.1
{[1.749079318, 1.749079318, 1.749079318, 1.749079318, 1.749079318, 1.749079318]}

    7.2
{[1.758855227, 1.758855227, 1.758855227, 1.758855227, 1.758855227, 1.758855227]}

    7.3
{[1.768534008, 1.768534008, 1.768534008, 1.768534008, 1.768534008, 1.768534008]}

    7.4
     $\emptyset$ 
    7.5
     $\emptyset$ 
    7.6
     $\emptyset$ 
    7.7
     $\emptyset$ 
    7.8
     $\emptyset$ 
    7.9
     $\emptyset$ 
    8.0
     $\emptyset$ 

```

(26)

> **for**  $a$  **from** 7.3 **by** 0.01 **to** 7.4

**do**

*print(a);*

$SEquP(GeneNet(0, a, 0.2, 2, m1, m2, m3, p1, p2, p3), [m1, m2, m3, p1, p2, p3]) ;$

**end do;**

7.3

{[1.768534008, 1.768534008, 1.768534008, 1.768534008, 1.768534008, 1.768534008]}

7.31

{[1.769496635, 1.769496635, 1.769496635, 1.769496635, 1.769496635, 1.769496635]}

7.32

{[1.770458315, 1.770458315, 1.770458315, 1.770458315, 1.770458315, 1.770458315]}

7.33

{[1.771419053, 1.771419053, 1.771419053, 1.771419053, 1.771419053, 1.771419053]}

7.34

{[1.772378849, 1.772378849, 1.772378849, 1.772378849, 1.772378849, 1.772378849]}

7.35

```

{[1.773337706, 1.773337706, 1.773337706, 1.773337706, 1.773337706, 1.773337706]}
    7.36
{[1.774295627, 1.774295627, 1.774295627, 1.774295627, 1.774295627, 1.774295627]}
    7.37
{[1.775252614, 1.775252614, 1.775252614, 1.775252614, 1.775252614, 1.775252614]}
    7.38
{[1.776208668, 1.776208668, 1.776208668, 1.776208668, 1.776208668, 1.776208668]}
    7.39
{[1.777163792, 1.777163792, 1.777163792, 1.777163792, 1.777163792, 1.777163792]}
    7.40
     $\emptyset$ 

```

(27)

```

>
> #9a,b)
> SEquP(ChemoStat(N, C, 2.5, 2.7), [N, C])
    {[5.083333333, 0.6666666667]} (28)
>
> #10)

```

```

> P := [[0.2, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8], [0.8/8, 0.2, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8], [0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8, 0.8/8], [0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8], [0.6/8, 0.4, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8], [0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.4, 0.6/8, 0.6/8, 0.6/8], [0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.6/8, 0.4, 0.6/8, 0.6/8, 0.4/8], [0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8], [0.4/8, 0.6, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.6, 0.4/8], [0.4/8, 0.4, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4, 0.6/8], [0.4/8, 0.4, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.4/8, 0.6]]]
```

```

P := [[0.2, 0.1000000000, 0.1000000000, 0.1000000000, 0.1000000000, 0.1000000000,
0.1000000000, 0.1000000000, 0.1000000000], [0.1000000000, 0.2, 0.1000000000,
0.1000000000, 0.1000000000, 0.1000000000, 0.1000000000, 0.1000000000],
[0.1000000000, 0.1000000000, 0.2, 0.1000000000, 0.1000000000, 0.1000000000,
0.1000000000, 0.1000000000, 0.1000000000], [0.07500000000, 0.07500000000,
0.07500000000, 0.4, 0.07500000000, 0.07500000000, 0.07500000000, 0.07500000000,
0.07500000000], [0.07500000000, 0.07500000000, 0.07500000000, 0.07500000000, 0.07500000000, 0.4,
0.07500000000, 0.07500000000, 0.07500000000, 0.07500000000], [0.07500000000,
0.07500000000, 0.07500000000, 0.07500000000, 0.4, 0.07500000000, 0.07500000000,
0.07500000000, 0.07500000000], [0.05000000000, 0.05000000000, 0.05000000000,
0.05000000000, 0.05000000000, 0.6, 0.05000000000, 0.05000000000], (29)

```

```
[0.050000000000, 0.050000000000, 0.050000000000, 0.050000000000, 0.050000000000,
0.050000000000, 0.050000000000, 0.6, 0.050000000000], [0.050000000000, 0.050000000000,
0.050000000000, 0.050000000000, 0.050000000000, 0.050000000000,
0.050000000000, 0.6]]
```

>  $\text{evalm}(P^{1000})$

$$\begin{aligned} & [[0.07692307655, 0.07692307655, 0.07692307655, 0.1025641021, 0.1025641021, \\ & \quad 0.1025641021, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307655, 0.07692307655, 0.07692307655, 0.1025641021, 0.1025641021, \\ & \quad 0.1025641021, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307655, 0.07692307655, 0.07692307655, 0.1025641021, 0.1025641021, \\ & \quad 0.1025641021, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307651, 0.07692307651, 0.07692307651, 0.1025641020, 0.1025641020, \\ & \quad 0.1025641020, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307651, 0.07692307651, 0.07692307651, 0.1025641020, 0.1025641020, \\ & \quad 0.1025641020, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307651, 0.07692307651, 0.07692307651, 0.1025641020, 0.1025641020, \\ & \quad 0.1025641020, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307651, 0.07692307651, 0.07692307651, 0.1025641020, 0.1025641020, \\ & \quad 0.1025641020, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307651, 0.07692307651, 0.07692307651, 0.1025641020, 0.1025641020, \\ & \quad 0.1025641020, 0.1538461530, 0.1538461530, 0.1538461530], \\ & \quad [0.07692307651, 0.07692307651, 0.07692307651, 0.1025641020, 0.1025641020, \\ & \quad 0.1025641020, 0.1538461530, 0.1538461530, 0.1538461530]] \end{aligned}$$

>

(30)