```
Laboratory 1: Modelling Change with Difference Equations
1. Write the first five terms of the following sequences:
   (a) a_{n+1} = 3a_n, a_0 = 1;
   (b) a_{n+1} = 3a_n (a_n + 1), a_0 = 0;
   (c) a_{n+1} = 2a_n^2, a_0 = -1;
  restart;
> n := 5;
                                                                                     (1)
                                      n := 5
Exercise (1.a)
> a[0] := 1;
                                                                                     (2)
                                      a_0 := 1
> for i from 1 to n do
    a[i] := 3 * a[i-1];
  end do:
> seq(a[i], i=0..n);
                                                                                     (3)
                                 1, 3, 9, 27, 81, 243
Exercise (1.b)
> a[0] := 0;
                                      a_0 := 0
                                                                                     (4)
> for i from 1 to n do
     a[i] := 3 * a[i-1] * (a[i-1] + 1);
  end do:
> seq(a[i], i=0..n);
                                   0, 0, 0, 0, 0, 0
                                                                                     (5)
Exercise (1.c)
> a[0] := -1;
                                     a_0 := -1
                                                                                     (6)
> for i from 1 to n do
     a[i] := 2 * a[i-1]^2;
  end do:
> seq(a[i], i=0..n);
```

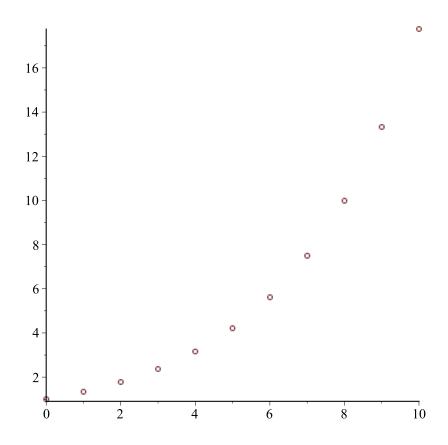
-1, 2, 8, 128, 32768, 2147483648

(7)

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2. Write the first ten terms of the sequence satisfying the following difference equations and
     draw the corresponding graph of the generated dynamical system:
     (a) \Delta a_n = \frac{1}{3}a_n, a_0 = 1;
     (b) \Delta a_n = 0.01 \cdot (200 - a_n), a_0 = 10;
     (c) \Delta a_n = 1.5 \cdot (100 - a_n), a_0 = 200;
> restart;
                                                         n := 10
                                                                                                                               (8)
Exercise (2.a)
> a[0] := 1;
                                                         a_0 := 1
                                                                                                                               (9)
> for i from 1 to n do
        a[i] := a[i-1] + 1/3 * a[i-1];
    end do:
> seq(a[i], i=0..n);
           1, \frac{4}{3}, \frac{16}{9}, \frac{64}{27}, \frac{256}{81}, \frac{1024}{243}, \frac{4096}{729}, \frac{16384}{2187}, \frac{65536}{6561}, \frac{262144}{19683}, \frac{1048576}{59049}
```

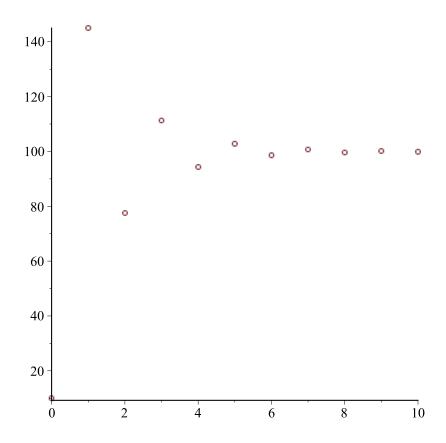
> plot([[j, a[j]]\$j=0..n], style=point, symbol=circle);

(10)



```
a_{10} := 28.16740574 (12)
```

```
> plot([[j, a[j]]$j=0..n], style=point, symbol=circle);
        28
                                                     o
        26
                                                 0
        24
        22
        20
                                  0
        18
        16
        14 -
                    0
        12 -
        10
                    2
                                       6
                                                          10
```



3. (Decay of Digoxin in the Bloodstream) Digoxin is used in treatment of heart disease. Doctors must prescribe an amount of medicine that keeps the concetration of digoxin in the bloodstream above an *effective level* without exceeding the *safe level*. For an initial dosage of 0.5 mg in the bloodstream, table shows the amount of digoxin a_n remaining in the bloodstream of a particular pacient after n days.

- (a) Get Δa_n and plot Δa_n versus a_n
- (b) Find a proportional constant for Δa_n and a_n .

```
Exercise (3.a)
> for i from 1 to 7 do
     delta a[i] := a[i+1] - a[i];
   end do:
> seq(delta_a[i], i=1..7);
                -0.155, -0.107, -0.074, -0.051, -0.035, -0.024, -0.017
                                                                                 (16)
> plot([[a[j], delta_a[j]]$j=1..7], style=point, symbol=circle,
   labels=[Digoxin, Change_of_digoxin], labeldirections=[HORIZONTAL,
   VERTICAL]);
                                 Digoxin
                            0.2
                                                         0.5
                  0.1
                                               0.4
                                     0.3
          -0.02
                    0
          -0.04
                         o
          -0.06
       Change_of_digoxin
          -0.08
          -0.10
                                          0
          -0.12
          -0.14
Exercise (3.b)
> k = (delta_a[6] - delta_a[4]) / (a[6] - a[4]);
                                k = -0.3139534884
                                                                                 (17)
```

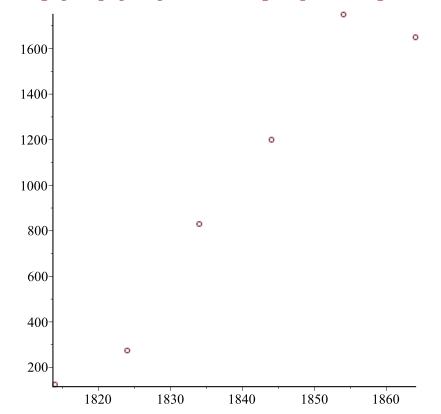
4. The following data were obtained for the growth of a sheep population introduced into new environment on the island of Tasmania:

```
Year 1814 1824 1834 1844 1854 1864
Population 125 275 830 1200 1750 1650
```

- (a) Plot the data. Is there a trend?
- (b) Formulate discrete dynamical system that resonable approximate the change observed. Compare real data with estimated data obtained from the model.

Exercise (4.a)

> plot([[y[j], p[j]]\$j=1..6], style=point, symbol=circle);



We cannot really notice a trend between the year and the population, we'll try to check for a trend between the population and delta population.

```
> for i from 1 to 5 do
    delta p[i] := p[i+1] - p[i];
  end do:
> seq(delta_p[i], i=1..5);
                             150, 555, 370, 550, -100
                                                                                (19)
> plot([[p[j], delta_p[j]]$j=1..5], style=point, symbol=circle,
  labels=[Population, Change of population], labeldirections=
  [HORIZONTAL, VERTICAL]);
                                          0
          500
          400-
      Change_of_population
          300
          200
          100
            0
                              800
                                   1000
              200
                   400
                         600
                                         1200
                                              1400
                                                   1600
                               Population
```

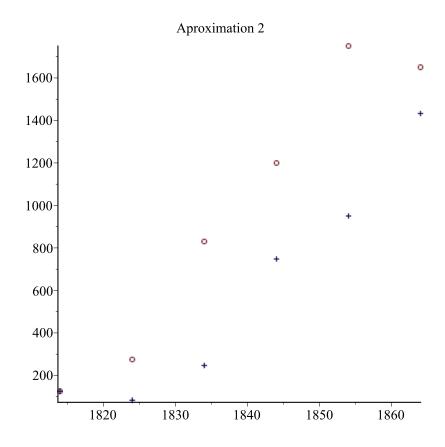
There seems to be a proportionality between populations 1, 3 and 4 and between populations 2, 3 and 5. We'll plot both to see which model is better.

0

Exercise (4.b)

-100

```
> seq(p_est[i], i=1..6);
                       125, \frac{6875}{37}, \frac{15125}{37}, \frac{45650}{37}, \frac{66000}{37}, \frac{96250}{37}
                                                                                             (21)
> est_data := [[y[j], p_est[j]]$j=1..6]:
> plot([real_data, est_data], style=point, symbol=[circle, cross],
   title="Aproximation \overline{1}");
                                Aproximation 1
          2500
          2000
                                                        å
          1500
                                              ŧ
          1000
           500
                         0
                   1820
                              1830
                                        1840
                                                   1850
                                                             1860
> k2 := (delta p[3] - delta p[2]) / (p[3] - p[2]);
                                        k2 := -\frac{1}{3}
                                                                                             (22)
> p est[1] := p[1]:
   for i from 2 to 6 do
      p_est[i] := p[i-1] + p_est[i-1] * k2;
> seq(p_est[i], i=1..6);
                       125, \frac{250}{3}, \frac{2225}{9}, \frac{20185}{27}, \frac{77015}{81}, \frac{348235}{243}
                                                                                             (23)
> est_data := [[y[j], p_est[j]]$j=1..6]:
> plot([real_data, est_data], style=point, symbol=[circle, cross],
    title="Aproximation 2");
```



It seems like the first model is a bit optimistic because k1 > 0, so it will always predict a larger value than the previous one, while model 2 is very pesimistic because k2 < 0, so it will always predict a smaller value than the previous one, which from the graph it's not available in most of the cases. Because we have a few available data points, it's hard to find a better model.