```
myModel[populationSize_, mutationRate_] := [
  (*Set initial condition*)
  abFrequency = 1; (*frequency of the allele to be followed (A) *)
  AbFrequency = 0;
  aBFrequency = 0;
  ABFrequency = 0;
  numberOfGenerations = 1000;
  (*Initialise list in which we will save the frequencies from every pokolenie*)
  abfrequencies = Table[0, {numberOfGenerations}];
  Abfrequencies = Table[0, {numberOfGenerations}];
  aBfrequencies = Table[0, {numberOfGenerations}];
  ABfrequencies = Table[0, {numberOfGenerations}];
  For [i = 1, i ≤ numberOfGenerations, i++,
    (*mutation*)
   abFrequency = (1 - mutationRate)^2 * abFrequency + (1 - mutationRate) *
       mutationRate * (AbFrequency + aBFrequency) + mutationRate<sup>2</sup> * ABFrequency;
   AbFrequency = (1 - mutationRate) * AbFrequency + (1 - mutationRate) *
       mutationRate * (ABFrequency + abFrequency) + mutationRate<sup>2</sup> * aBFrequency;
   aBFrequency = (1 - mutationRate) * aBFrequency + (1 - mutationRate) *
       mutationRate * (ABFrequency + abFrequency) + mutationRate<sup>2</sup> * AbFrequency;
   ABFrequency = (1 - mutationRate) * ABFrequency + (1 - mutationRate) *
       mutationRate * (aBFrequency + AbFrequency) + mutationRate<sup>2</sup> * abFrequency;
    (*selection*)
   abw = 1;
   Abw = 1.01;
   aBw = 1.01;
   ABw = 1.012;
   w = abw * abFrequency + Abw * AbFrequency + aBw * aBFrequency + ABw * ABFrequency;
   abFrequency = abFrequency * \frac{abw}{...};
   AbFrequency = AbFrequency * \frac{Abw}{...};
   aBFrequency = aBFrequency * \frac{aBw}{};
   ABFrequency = ABFrequency * \frac{ABw}{};
    (*crossing*)
   De = abFrequency * ABFrequency - AbFrequency;
   r = 0.1;
   abFrequency = abFrequency - r * De;
   AbFrequency = AbFrequency + r * De;
   aBFrequency = aBFrequency + r * De;
   ABFrequency = ABFrequency - r * De;
    (*sampling*)
```

```
abNa = RandomVariate[BinomialDistribution[populationSize, abFrequency]];
   abFrequency =
                  populationSize
   AbNa = RandomVariate[BinomialDistribution[populationSize, AbFrequency]];
                        AbNa
   AbFrequency =
                  populationSize
   aBNa = RandomVariate[BinomialDistribution[populationSize, aBFrequency]];
                        aBNa
   aBFrequency = -
                  populationSize
   ABNa = RandomVariate[BinomialDistribution[populationSize, ABFrequency]];
                        ABNa
   ABFrequency =
                  populationSize
   (*saving values*)
   abfrequencies[[i]] = abFrequency;
   Abfrequencies[[i]] = AbFrequency;
   aBfrequencies[[i]] = aBFrequency;
   ABfrequencies[[i]] = ABFrequency;
  ];
  Show [
   ListPlot[abfrequencies, PlotRange → Automatic, PlotStyle → Gray],
   ListPlot[Abfrequencies, PlotRange → Automatic, PlotStyle → Red],
   ListPlot[aBfrequencies, PlotRange → Automatic, PlotStyle → Green],
   \textbf{ListPlot}[\texttt{ABfrequencies, PlotRange} \rightarrow \texttt{Automatic, PlotStyle} \rightarrow \texttt{Blue}]
myModel[100000, 0.0001]
1.0
8.0
0.6
0.4
0.2
                      400
                                600
                                           800
                                                     1000
```