3.d term

Population genetics

Genetic drift and mutation

Problems

A p: 0'875

a q: 0'125

4 individuals -> 8 alleles

No a >8A > 0'875 **8 = 0'3 44 > probability of no a (P(A) = P(noa))

NoA > 8a > 0'125 ** = 5'96 · 10-8 -> probability of no A (P(A) = P(noa))

P2+2pq+q2=1 > 2pq=1 ptq2 > 2pq = 1-(01252+08752) = d219

P(Ama) P(Ana) = 1-P(A)-P(a) -> P(Ana)=1-{10'344-5'96.10-8}=0'656

A a \Rightarrow 12 individuals \Rightarrow 24 alleles

40% heterozygous \Rightarrow num of generalions \Rightarrow Ht = 1-09=09 in 10% heterozygous

H_t = Ho $\left(1 - \frac{1}{2N}\right)^{\frac{1}{2}} \Rightarrow \ln\left(\frac{1+1}{2N}\right) \Rightarrow + \frac{\ln\left(\frac{1+1}{2N}\right)}{\ln\left(1 - \frac{1}{2N}\right)} \Rightarrow + \frac{\ln\left(\frac{1+1}{2N}\right)}{\ln\left(\frac{1+1}{2N}\right)} \Rightarrow + \frac{\ln\left(\frac{1+1}{2N}\right)}$

N=95I newallele $Po \rightarrow \frac{1}{2N} = Po \Rightarrow \frac{1}{24S} = 0'0053 \Rightarrow t$ here is a 0'53% phrobability that it will drift to fixation $Plost = 1 - Po \Rightarrow Plost = 1 - 0'0053 = 0'995 \Rightarrow t$ here is a 99'5% probability of loss $T_{Cix} = -4N = \frac{(1-P)h(1-P)}{P} = -4'95 = -4'95 = 378'9 \Rightarrow 379 \Rightarrow 3$

P1=0175 N= 40

P4 = 0'75

P(kallelesin nalleles)= 1! (n-k) pk, qn-k

K=2N. pi + K=2. 40. 0975= 60

9=1-1-1- 9=1-075=075

1=2N=) 1=2-40=80

P(60 in 80) = 80) (80-60) . P01750. 0'25 (80-60) = 0'10 25

There is a 10'25% chance that inthenext generation the allele A frequency is of 0'75

N=12 1 36 F

A = 0'5 = Po

a = 0's = 90 P1 = 0'48 91 = 1-04588 = 0'642

P2 = 0'583 9 = 1- d 583 = 0'417

k= 2N. Po > k= 12.0'5 = 6

1=2N= 2.6=12

 $P_1 = \frac{12!}{6!(12-6)!} \cdot 0458^6 \cdot 0542^{(12-6)} = 02162$

 $P_z = 4\frac{12!}{6!(126)!}$. $6'583^6$. $0'417^{(12-6)} = 0'19077$

Pd = 0'375 9= 1-0'375=0'625 kd=18-0'375=4'5~5

P= 12! 10'375 . 0'625 = 0'22

P(0375) in PrandP2) = 022.0'22=004787 4'787% of that second generation

6 p1s Genetic drift theory

Po=0'5 p90=0's = 2A Za

+=2

P2=0'25 92=0'75 > 1A 3a

P(0'25|0'5) = P(1A | ZA) = P(1allele Aingen Z) = 1200'4244

4 > table 0 1 Z 3 4 negen

O'ores

O'allele 1 0 0'25 0'0440 0

ingen 1 0'355
0'0625

P(1all in gen 2) = (0.0'0625) + (0'4219.0'25)+(0'250'375)+(0'049.0'25)
+ (0.0'0625) = 0 + 0'1055 + 0'0938 + 0'01173 + 0 = 0'21103

The probability that a population of zindividuals (4 alleles)
that start with p=0'5 and have p=0'25 in 2 generations is
of 0'21103

NET = \$5000 = 1100 males

NE = \$5000 - 1100 = \$3400 females Sadultidividuals allin groups of \$0 to 1M

Nes Not 100 - 53400 = 4312

Non the Moots9900

Assuming equal probability of M/F individual indescendency

27500 = M 27500 = F (there will be males without a group)

Mutation

A → a

µ: 500002

Po=015

Pro? Pt = Po (1-y)[†]

Pro = o's (1-0'00002)¹⁰ = 0'4999 00009 = Frequery of Air 10 generations

Pro = o's (1-0'00002)¹⁰⁰ = 0'49900099 = Frequery of Air 100 generations

Pro = o's (1-0'00002)¹⁰⁰⁰ = 0'49009 = Frequery of Air 1000 generations

Pro = o's (1-0'00002)¹⁰⁰⁰ = o'4009 = Frequery of Air 10000 generations

Pro = o's (1-0'00002)¹⁰⁰⁰⁰ = o'409365 = Frequery of Air 100000 generations

Its a very slow change inallele frequery

 $P_{s}=o'8 \qquad q_{s}=o'2$ $P_{t}=\frac{v}{\mu+v}+\left(P_{0}-\frac{v}{\mu+v}\right)\left(1-\mu-v\right)^{+}$ $q_{1}=\frac{o'oods}{o'oos+o'oods}+\left(o'z-\frac{o'oods}{o'oos+o'oods}\right)\left(1-o'oos-o'oods\right)^{1}=o'2002$ In the First generation q=o'2002 $q_{2}=\frac{o'oods}{o'oos+o'oods}+\left(o'z-\frac{o'oods}{o'oos+o'oods}\right)\left(1-o'oos-o'oods\right)^{2}=o'200348$ In the second generation q=200348 $q_{3}=\frac{o'oods}{o'oos+o'oods}+\left(o'z-\frac{b'o'oods}{b'oos+o'oods}\right)\left(1-o'oos-o'oods\right)^{3}=o'2005961$ In the third generation q=0'2005961

 $P = \frac{V}{\mu + V} \rightarrow P = \frac{0'00AS}{0'005t0'001S} = 0'2308$ $q = 1 - p \Rightarrow q = 1 - 0'2308 = 0'7692$

Allele Frequencies in equilibriumare 9=07692 p=02388