

Power Analysis

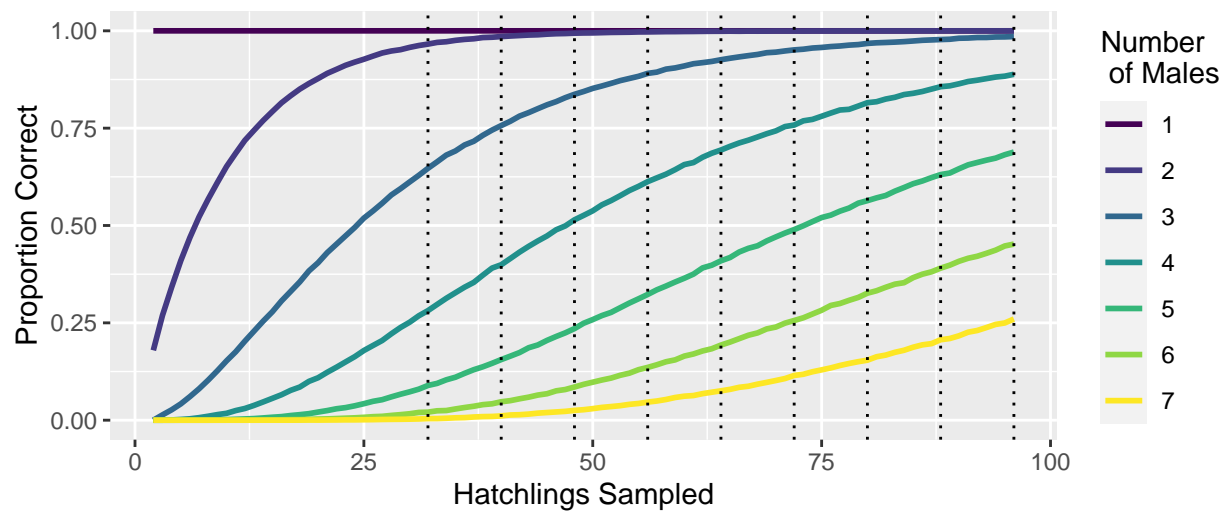
Question 1: How many hatchlings should be sampled from a nest to robustly estimate the number of males that contributed to it?

Assuming one dominant sire that fertilizes 90% of eggs

```
source('hatchlings_to_sample.R')

hatchlings_to_sample(n_hatchlings = 100,
  max_hatchlings = 96,
  max_males = 7,
  breeding = 'dominant',
  n_sims = n_sims,
  dom = 0.9,
  n_sizes = c(32, 40, 48, 56, 64, 72, 80, 88, 96))
```

[[1]]



##

[[2]]

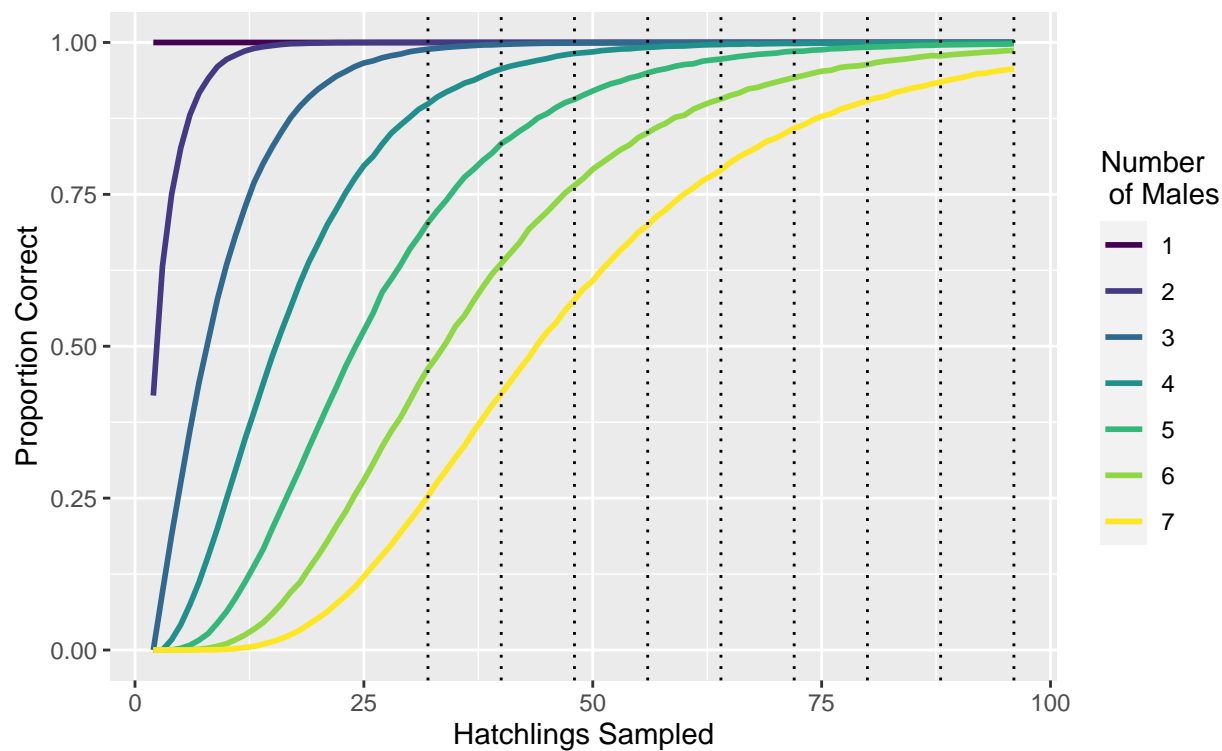
##	Males	32	40	48	56	64	72	80	88	96
## 1	1	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000
## 2	2	0.96589	0.98519	0.99398	0.99717	0.99891	0.99951	0.99977	0.99988	0.99999
## 3	3	0.64689	0.75664	0.83692	0.89070	0.92573	0.95104	0.96769	0.97759	0.98529
## 4	4	0.28144	0.39921	0.51453	0.61268	0.69384	0.75795	0.81543	0.85653	0.88835
## 5	5	0.08918	0.15564	0.23521	0.32282	0.40934	0.48947	0.56385	0.63079	0.68862
## 6	6	0.02068	0.04732	0.08516	0.13555	0.19292	0.25597	0.32618	0.39050	0.45227
## 7	7	0.00412	0.01094	0.02521	0.04653	0.07537	0.11482	0.15412	0.20662	0.25980

Assuming one dominant sire that fertilizes 70% of eggs

```
source('hatchlings_to_sample.R')

hatchlings_to_sample(n_hatchlings = 100,
  max_hatchlings = 96,
  max_males = 7,
  breeding = 'dominant',
  n_sims = n_sims,
  dom = 0.7,
  n_sizes = c(32, 40, 48, 56, 64, 72, 80, 88, 96))
```

[[1]]



##

[[2]]

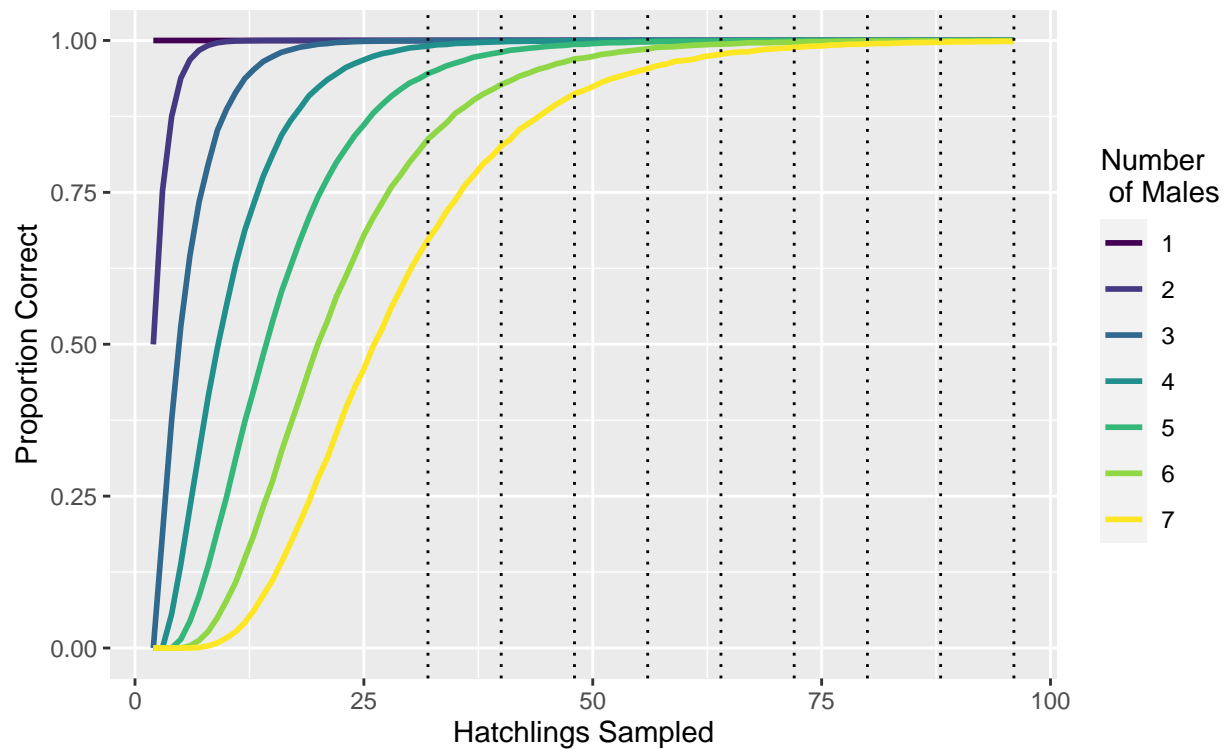
##	Males	32	40	48	56	64	72	80	88	96
## 1	1	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000
## 2	2	0.99996	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000
## 3	3	0.98895	0.99680	0.99910	0.99979	0.99995	0.99998	0.99998	0.99999	1.00000
## 4	4	0.89881	0.95657	0.98197	0.99184	0.99661	0.99824	0.99933	0.99962	0.99984
## 5	5	0.70341	0.83342	0.90640	0.94998	0.97269	0.98555	0.99220	0.99595	0.99791
## 6	6	0.46308	0.63617	0.76457	0.85138	0.90689	0.94222	0.96380	0.97798	0.98680
## 7	7	0.25325	0.42173	0.57643	0.69932	0.78982	0.85888	0.90446	0.93467	0.95643

Assuming one dominant sire that fertilizes 50% of eggs

```
source('hatchlings_to_sample.R')

hatchlings_to_sample(n_hatchlings = 100,
  max_hatchlings = 96,
  max_males = 7,
  breeding = 'dominant',
  n_sims = n_sims,
  dom = 0.5,
  n_sizes = c(32, 40, 48, 56, 64, 72, 80, 88, 96))
```

[[1]]



##

[[2]]

##	Males	32	40	48	56	64	72	80	88	96
## 1	1	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000
## 2	2	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000
## 3	3	0.99982	0.99999	0.99999	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000
## 4	4	0.99080	0.99785	0.99954	0.99985	0.99996	1.00000	1.00000	1.00000	1.00000
## 5	5	0.94553	0.98070	0.99377	0.99764	0.99933	0.99975	0.99988	0.99994	0.99999
## 6	6	0.83742	0.92746	0.96952	0.98603	0.99395	0.99747	0.99907	0.99948	0.99983
## 7	7	0.67225	0.82614	0.91243	0.95360	0.97722	0.98841	0.99437	0.99706	0.99832