

A Population Genetic Report

using PopGenReport Ver. 1.6.1 .

Adamack & Gruber (2013)

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Contents

1 Counts

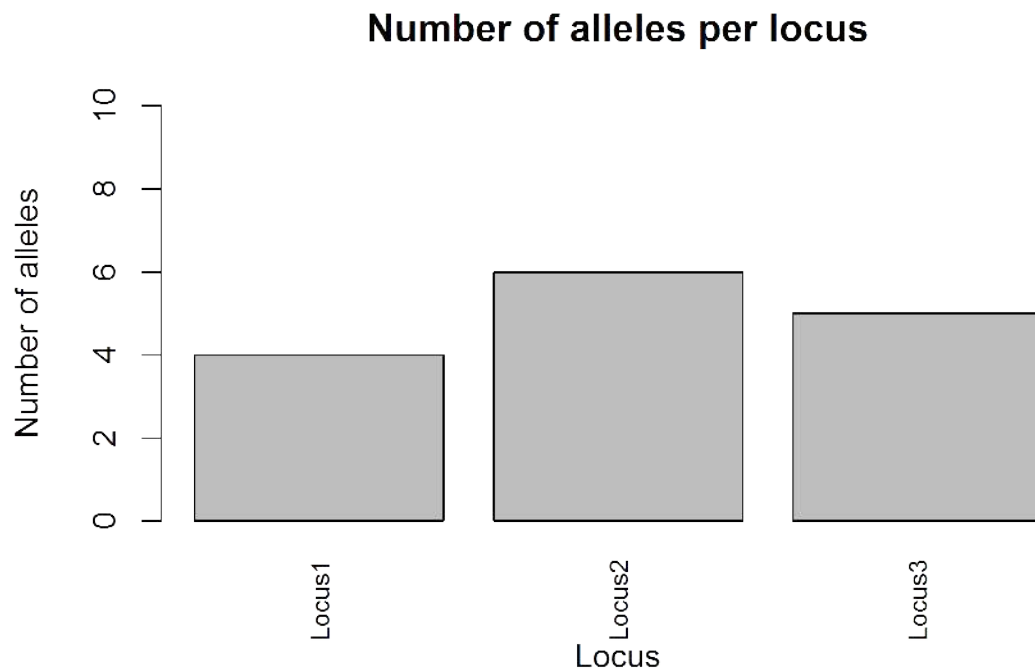
2

1 Counts

This analysis looks at 71 individuals.

The mean number of alleles per locus (across all locations): 5

The percentage of missing data was 0%.



The individuals were sampled from the following locations in the following numbers:

| population | ACT | NSW | QLD |
|------------|-----|-----|-----|
| # ind | 18 | 30 | 23 |

Table 1: Number of individuals per population

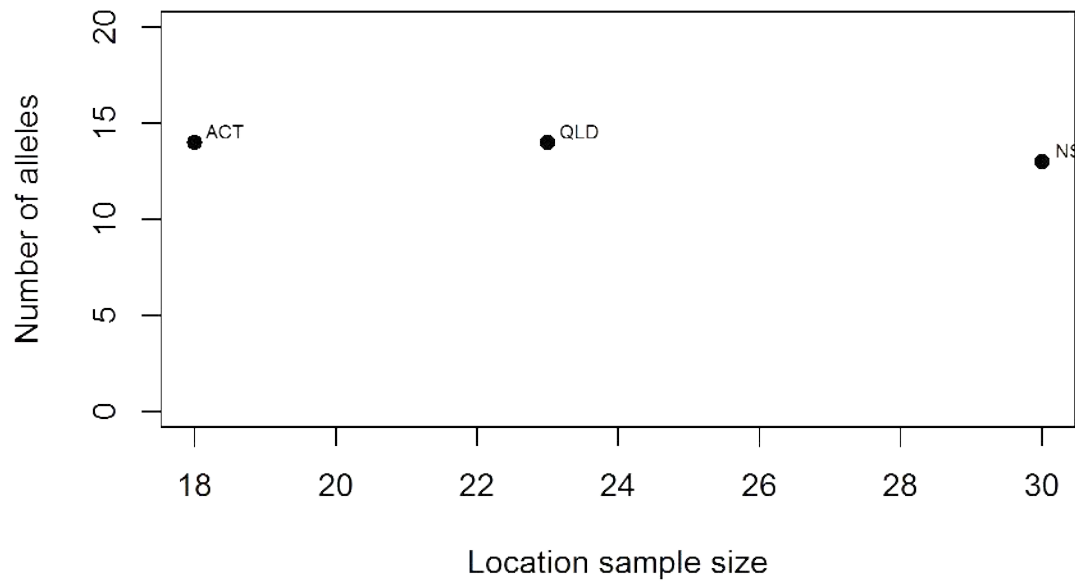
The total number of alleles sampled across all locations was 15

The total number of alleles seen in each sampling location was:

| population | ACT | NSW | QLD |
|------------|-----|-----|-----|
| # alleles | 14 | 13 | 14 |

Table 2: Number of alleles per population

Number of alleles vs location sample size



The number of alleles per locus (across all subpopulations):

| locus | Locus1 | Locus2 | Locus3 |
|-----------|--------|--------|--------|
| # alleles | 4 | 6 | 5 |

Table 3: Number of alleles per locus across all subpopulations