Supporting Information

**Strategies for determining kinship in wild populations using genetic data**

*Ecology and Evolution*

**Table S1** Numbers of microsatellites (STRs) and single-nucleotide polymorphisms (SNPs)

with equal power for kinship analyses

|  |  |  |
| --- | --- | --- |
| Number of STRs | Number of SNPs | Reference |
| 1 | 10 | Wang & Santure 2009 |
| 6 | 32 | Glaubitz *et al.* 2003 |
| 10 | 43 | Herráez *et al.* 2005 |
| 10 | 60 | Rohrer *et al.* 2007 |
| 11 | 80 | Hauser *et al.* 2011 |
| 13 | 63 | Sellars *et al.* 2014 |
| 13 | 59 | Sellars *et al.* 2014 |
| 14 | 100 | Morin *et al.* 2004 |
| 15 | 102 | Weinman *et al.* 2015 |
| 16 | 106 | Telfer *et al.* 2015 |
| 17 | 95 | Steele *et al.* 2013 |
| 18 | 100 | Glaubitz *et al.* 2003 |

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