**Appendix S1.**

# Johannesen J, FabritzekAG, Ebner B, BikarS-E. Characterisation of microsatellite and SNP markers from Miseq and genotyping-by-sequencing data among parapatric *Urophora cardui* (Tephritidae) populations

**Barcode adapters,** restriction enzyme EcoR1: 5- G’AATT’C -3

|  |  |  |  |
| --- | --- | --- | --- |
| Name | Barcode | BCN | Sequence **5‘ –> 3‘** |
| **BC-18-EcoR1** | **taggaa** | 6 | **AATTttccta**AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT |
|  |  |  |  |
| **BC-20-EcoR1** | **cttgctt** | 7 | **AATTaagcaag**AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT |
|  |  |  |  |
| **BC-24-EcoR1** | **tagcatgc** | 8 | **AATTgcatgcta**AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT |
|  |  |  |  |
| **BC-25-EcoR1** | **tgcaagga** | 8 | **AATTtccttgca**AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT |
|  |  |  |  |
| **BC-26-EcoR1** | **aaccgag** | 7 | **AATTctcggtt**AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT |
|  |  |  |  |
| **BC-JJ1-EcoR1** | **ttgcga** | 6 | **AATTtcgcaa**AGATCGGAAGAGCGTCGTGTAGGGAAAGAGTGT |
|  |  |  |  |

**Common adapters 5‘ –> 3‘**

|  |  |
| --- | --- |
| Common top**Com-Top- EcoR1** | **AATT**AGATCGGAAGAGCACACGTCTGAACTCCAGTCA |
| Common bottom**Com-Bottom** | TGACTGGAGTTCAGACGTGTGCTCTTCCGATCT |

**PCR primers with Index ID barcodes** (underlined = Barcode (1) & common bottom (2) sequences)

|  |  |
| --- | --- |
| Primer | Sequence 5’ –> 3’ |
| PCR Primer 1 **GBS P1 univ** | AATGATACGGCGACCACCGAGATCTACACTCTTTCCCTACACGACGCTCTTCCGATCT |
|  |  |
| PCR Primer 2 |  |
| **TS08** (acttga) | CAAGCAGAAGACGGCATACGAGAT**tcaagt**GTGACTGGAGTTCAGACGTGTGCTCTTCCGATC |
| **TS13** (agtcaa) | CAAGCAGAAGACGGCATACGAGAT**ttgact**GTGACTGGAGTTCAGACGTGTGCTCTTCCGATC |
| **TS14** (agttcc) | CAAGCAGAAGACGGCATACGAGAT**ggaact**GTGACTGGAGTTCAGACGTGTGCTCTTCCGATC |
| **TS15** (atgtca) | CAAGCAGAAGACGGCATACGAGAT**tgacat**GTGACTGGAGTTCAGACGTGTGCTCTTCCGATC |
| **TS16** (ccgtcc) | CAAGCAGAAGACGGCATACGAGAT**ggacgg**GTGACTGGAGTTCAGACGTGTGCTCTTCCGATC |