

## PCR PROTOCOL

**Purpose:** trnH-psbA DNA barcoding of plants

**Original Source:** Sang et al. 1997; Tate and Simpson 2003

**Document By:** BG (1/06/2020).

**Notes:** Protocols developed/implemented by Smithsonian/Guelph. Targets a non-coding/length-variable locus.

### Primers:

|          |                        |
|----------|------------------------|
| psbA3_f  | GTTATGCATGAACGTAATGCTC |
| trnHf_05 | CGCGCATGGTGGATTCAATCC  |

### Reaction (25 µL)

| REAGENTS (Final concentration)      | 1 X (µL) |
|-------------------------------------|----------|
| DMSO (3%)                           | 0.3      |
| MOLECULAR WATER                     | 6.320    |
| 5X HF Buffer (w MgCl <sub>2</sub> ) | 2        |
| 10 mM dNTPs                         | 0.056    |
| PRIMER1 – 10µM (0.2µM final)        | 0.1      |
| PRIMER2 – 10µM (0.2µM final)        | 0.1      |
| Phusion Hot Start II DNA Polymerase | 0.125    |
| <b>DNA TEMPLATE</b>                 | <b>1</b> |

**\*Make extra master mix and include a negative and positive control.**

**Program:** Phusion Trnh-Psba Bar on Eppendorf Mastercycler Nexus GSX1

| Process               | Temperature (°C) | Time   |
|-----------------------|------------------|--------|
| Initial denaturing    | 98               | 45 s   |
| 35 cycles: Denaturing | 98               | 10 s   |
| 35 cycles: Annealing  | 64               | 30 s   |
| 35 cycles: Extension  | 72               | 40 s   |
| Final Extension       | 72               | 10 min |
| Hold                  | 4                | Hold   |

Approximate total time: 80 min.