
The eINTACT method for studying nuclear changes in host plant cells targeted by bacterial effectors in native infection contexts

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1 **This PDF file includes:**

2 **Supplementary Tables 1 and 2**

3 **Supplementary Table 1.** Sequences of oligonucleotide primers for the semi-quantitative RT-
4 PCR analysis.

| Gene/Fragment | Oligo | Sequence (5' -> 3') ^a |
|---------------|---------|-------------------------------------|
| <i>NTF</i> | forward | ATGAATCATTCAGCGAAAACCA |
| | reverse | AGCAGCAGCAGCAGCCTTGTACAGCTCGTCCATGC |
| <i>TUB2</i> | forward | GAGCCTTACAACGCTACTCTGTCTGTC |
| | reverse | ACACCAGACATAGTAGCAGAAATCAAG |

5 ^a Oligonucleotide primers are synthesised at Eurofins Genomics.

6 **Supplementary Table 2.** Plasmids used in this protocol.

| Plasmid | Description (Resistance) | References |
|---------|---|------------|
| pYY1704 | <i>pBs3:RedNTF:tNOS</i> (spectinomycin) | 12 |
| pDP085 | <i>pUBQ10:BirA</i> (spectinomycin) | 23 |
| pDSK | A vector for expressing bacterial genes after a <i>lpp</i> promoter (spectinomycin) | 37, 40 |
| pDS300F | <i>plpp:AvrBs3</i> (spectinomycin) | 37, 40 |

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