

A simple and fast triplex-PCR for the identification of milk's animal origin in Halloumi cheese and yoghurt

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SUPPLEMENTARY FILE

Table 1: Origin of primers, gene sequence position from start site, nucleotide sequence, melting temperature (T_m), and size of product expected from PCR.

| Origin | Gene Sequence Position | Primer Nucleotide Sequence | Amplicon , bp | T_m , °C |
|-------------------------------------|------------------------|--------------------------------------|---------------|------------|
| Universal, Forward (UN-FW) | 400 | 5' TGAGGACAAATATCATTYTGAGGRGC 3' | - | 61.6 |
| <i>Bos taurus</i> , Reverse (BO-RV) | 666 | 5' TAAGATGTCCTTAATGGTATAGTAG 3' | 287 | 56.4 |
| <i>Capra hircus</i> Reverse (CA-RV) | 693 | 5' TTAGAACAAGAATTAGTAGCATGGCG 3' | 313 | 60.1 |
| <i>Ovis aries</i> Reverse (OV-RV) | 710 | 5' GCGTGAATAGTACTAGTAGCATGAGGATGA 3' | 336 | 66.8 |

^aThe UN12S pair of primers were designed to bind and amplify the same complementary DNA sequence of the 12S rRNA gene in all mammalian species (Tobe and Linacre, 2008b)

Table 2. DNA amounts extracted from commercially available Cyprus Halloumi and Yoghurt products

| Halloumi ID code* | Milk content Species** | A260/A280 | Total DNA, ng | Yogurt ID code* | Milk content Species** | A260/A280 | Total DNA, ng |
|-------------------|--------------------------|-----------|---------------|-----------------|------------------------|-----------|---------------|
| H1 | 100% goat | 1.88 | 540 | Y1 | 50: 50% Sheep: goat | 1.94 | 6.66 |
| H2 | 100% goat | 1.66 | 675 | Y2 | 100% cow | 1.82 | 1.10 |
| H3 | 100% goat | 1.87 | 525 | Y3 | 100% cow | 1.94 | 10.05 |
| H4 | 50: 50, % Sheep: goat | 1.86 | 555 | Y4 | 100% sheep | 1.67 | 1.85 |
| H5 | 50: 50, % Sheep: goat | 1.75 | 630 | | | | |
| H6 | 50: 50, % Sheep: goat | 1.84 | 510 | | | | |

*The Halloumi and yogurt samples obtained from supermarkets were coded to protect the anonymity of the producer;

**The species origin and % milk content are as specified in the marketing labels of each product.