**Table 2**. ISSR primer core sequences, number of bands and polymorphism content

for *Tulipa suaveolens*

|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| **Primer code** | **Primer sequence** | **PB** | **PIC** | **RP** | **MRP** | **MI** | **H’** |
| UBC810 | (GA)8T | 25 | 0.26 | 14.4 | 0.57 | 7.4 | 0.404 |
| UBC811 | (GA)8C | 34 | 0.21 | 16.2 | 0.48 | 8.5 | 0.350 |
| UBC816 | (CA)8T | 15 | 0.18 | 4.44 | 0.30 | 3.1 | 0.294 |
| UBC824 | (TC)8G | 29 | 0.24 | 15.8 | 0.55 | 7.9 | 0.368 |
| UBC827 | (AC)8G | 30 | 0.27 | 18.6 | 0.62 | 9.4 | 0.409 |
| UBC836 | (AG)8YT | 30 | 0.27 | 18.7 | 0.63 | 9.5 | 0.417 |
| UBC841 | (GA)8YT | 17 | 0.37 | 13.6 | 0.80 | 7.3 | 0.545 |
| UBC843 | (CT)8RA | 24 | 0.26 | 16.5 | 0.69 | 7.2 | 0.399 |
| UBC845 | (CT)8RG | 32 | 0.30 | 22.9 | 0.72 | 11.3 | 0.468 |
| UBC851 | (GT)8YG | 14 | 0.33 | 11.6 | 0.83 | 5.4 | 0.492 |
| **Average** |  | **25** | **0.27** | **15.3** | **0.62** | **7.7** | **0.415** |

PB − No. of polymorphic bands, PIC − polymorphic information content, RP − Resolving power, MRP − Mean resolving power, MI – the Marker Index, H’ – the Shannon’s diversity index