Correlation coefficients for AUC-based MS-HRM calibration curves and counted methylation levels

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| --- | --- | --- | --- | --- |
| Locus name | Primer set | R2 | Methylation [%](n = 10) | ± SD |
| M | M HRM | 0.957 | 95.43 | 5.85 |
| M | M HRM Wojdacz | 0.846 | 148.58 | 8.68 |
| IM | IM HRM | 0.930 | 16.73 | 3.72 |
| U | U HRM | 0.852 | 1.95 | 0.69 |
| U | U HRM Wojdacz | 0.952 | 4.88 | 7.82 |

M - methylated locus, IM – intermediately methylated locus, U – unmethylated locus, R2 – square of the correlation coefficient, SD – standard deviation