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| --- | --- | --- | --- | --- | --- |
| Table 3. Percentage (%) of methylated CpGs across the entire amplicon. | | | | | |
| Gene name | **Amplicon** | **RWPE-1** | **CTPE** | **CAsE-PE** | **B26** |
| *S100P* | Amplicon #1 | 42.0 | 25.7 | 22.5 | 42.2 |
| *S100P* | Amplicon #2 | 71.7 | 43.9 | 29.2\* | 90.0 |
| *HYAL1* | Amplicon #1 | 100 | 71.9\* | 71.9\* | 97.2 |
| *HYAL1* | Amplicon #2 | 92.9 | 15.4\* | 7.6\* | 97.1 |
| *NTM* | Amplicon #1 | 1.4 | 19.0\* | 25.5\* | 2.4 |
| *NTM* | Amplicon #2 | 0.4 | 29.4\* | 5.7 | 0.6 |
| *NES* | Amplicon #2 | 68.6 | 88.6 | 75.4 | 84.1\* |
| *ALDH1A1* | Amplicon #1 | 3.3 | 11.3 | 7.4 | 5.0 |

The percentage of methylated CpGs in each clone were compared by Kruskal-Wallis test and \* *p* <0.05 relative to RWPE-1 by Dunn’s post-hoc analysis.