Accession Number: 18–341–140130 Report Date: 2018–12–16 Run Name: MGMT\_1962

Assay: MGMT/SAR Methylation Result: Fail Total % Methylation: 18%

MGMT CpG Sites 77 & 78

25% Methylated

2570

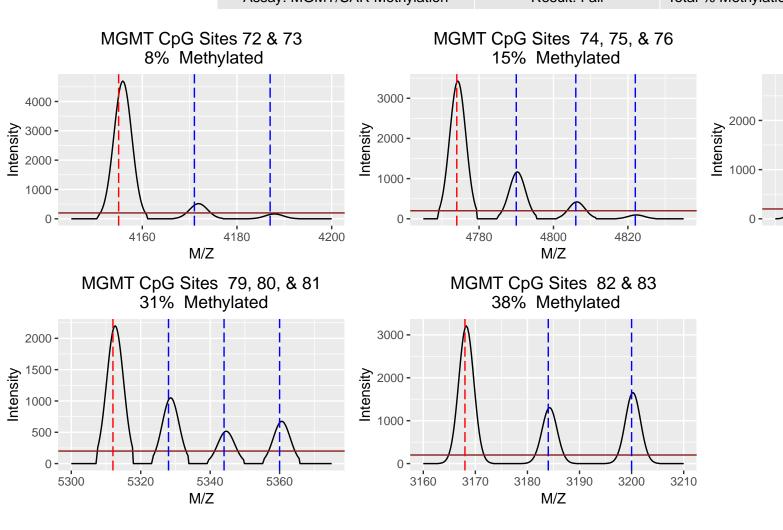
M/Z

2560

2550

2590

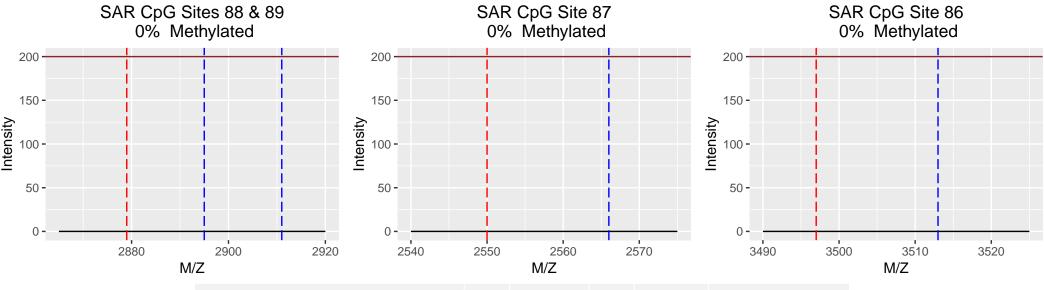
2580



|                                     |       |       |       |       |       | Pass/Fail | QC Check            |
|-------------------------------------|-------|-------|-------|-------|-------|-----------|---------------------|
| Np mean                             |       |       | 49.25 |       |       | Pass      | >=10                |
| Number of spectra                   |       |       | 5/5   |       |       | Pass      | >=3/5               |
| Warning flags                       |       |       | ok    |       |       | Pass      | ok                  |
| CpG unit # (CpGs/unit)              | 1 (2) | 2 (3) | 3 (2) | 4 (3) | 5 (2) |           |                     |
| Sum of peak(s) intensity per unit   | 5378  | 5112  | 4471  | 4441  | 6173  | Pass      | >=600 for each unit |
| Methylation value per unit (%)      | 8%    | 15%   | 25%   | 31%   | 38%   | Pass      | Calculated          |
| Total methylation from MGMT and SAR |       |       | 18%   |       |       | Pass      | Calculated          |

Accession Number: 18–341–140130 Report Date: 2018–12–16 Run Name: MGMT\_1962

Assay: MGMT/SAR Methylation Result: Fail Total % Methylation: 18%



|                                     |       |            |       | Pass/Fail | QC Check            |
|-------------------------------------|-------|------------|-------|-----------|---------------------|
| Np mean                             |       | 0          |       | Fail      | >=10                |
| Number of spectra                   |       | 0/5        |       | Fail      | >=3/5               |
| Warning flags                       |       | bad recal. |       | Fail      | ok                  |
| CpG unit # (CpGs/unit)              | 1 (2) | 2 (1)      | 3 (1) |           |                     |
| Sum of peak(s) intensity per unit   | 0     | 0          | 0     | Fail      | >=600 for each unit |
| Methylation value per unit (%)      | 0%    | 0%         | 0%    | Review    | Calculated          |
| Total methylation from MGMT and SAR |       | 18%        |       | Pass      | Calculated          |

| If                               | Then   |
|----------------------------------|--|
| Total % methylation is 0 – 9%    | The sample is reported as NOT DETECTED for MGMT promoter methlyation |
| Total % methylation is 10 – 29%  | The sample is reported as LOW LEVEL for MGMT promoter methlyation    |
| Total % methylation is 30 – 100% | The sample is reported as DETECTED for MGMT promoter methlyation     |

| Control                 | Percent Methylation | Pass/Fail |
|-------------------------|---------------------|-----------|
| Non-methylated Control  | 0–9%                | Pass      |
| 20% Methylated Control  | 10–40%              | Pass      |
| 100% Methylated Control | 90–100%             | Pass      |
| NTC                     | np mean <10         | Pass      |