



## Methylation Services Order

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To be completed by the user

Requested by: \_\_\_\_\_

PO #: \_\_\_\_\_

Bill to: \_\_\_\_\_

Phone #: \_\_\_\_\_

Institution: \_\_\_\_\_

Date: \_\_\_\_/\_\_\_\_/\_\_\_\_

Email: \_\_\_\_\_

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### Targeted - Sequenom EpiTyper:

Number of CpG sites requested: \_\_\_\_\_

Number of amplicons (primer sets): \_\_\_\_\_

Number of samples assayed: \_\_\_\_\_ (minimum recommended >30)

Bisulfite treatment required: \_\_\_\_\_ YES \_\_\_\_\_ NO

Gene\_CpG identifiers, sequences and 96-W DNA plate IDs must be emailed to [Michael.Bianchi@RoswellPark.org](mailto:Michael.Bianchi@RoswellPark.org)

### Whole Genome - Illumina Human Methylation 27:

Number of samples assayed: \_\_\_\_\_ (minimum = 12)

Bisulfite treatment required: \_\_\_\_\_ YES \_\_\_\_\_ NO

User Signature \_\_\_\_\_

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MICROARRAY FACILITY use only

Assay ID #: \_\_\_\_\_

Date Rec'd: \_\_\_\_/\_\_\_\_/\_\_\_\_

Rec'd by: \_\_\_\_\_

RPCI DNA Microarray Facility  
 BLSC, Room L1-130  
 Buffalo, NY 14263 USA  
 Telephone: 716 845-3361  
 Fax: 716 845-1579