sequencingcenter

John Wayne Cancer Institute
Providence Saint John's Health Center

METHYLATION ARRAY (EPIC) SAMPLE SUBMISSION FORM

INSTRUCTIONS: Please complete and email this form along with the MethylationEPIC Sample Sheet to hsut@jwci.org prior to sample submission.

REQUESTOR INFORMATION

Institution:	
Date:	
Requestor Name:	
Phone:	
Email:	
Principal Investigator:	
Phone:	
Email:	
SAMPLE INFORMATION	
Total Number of Samples:	
Sample Origin- check all that apply:	☐ Human Cell Lines☐ Human Tumor Tissue (Fresh or Frozen)☐ Human FFPE Tissue
DNA Quantification Method:	☐ PicoGreen dsDNA☐ Other—Specify method(s) on Sample Sheet

DATA ANALYSIS		
Would you like data analysis to be conducted?	□ YES	
	□ NO	
RESIDUAL MATERIALS		
□ DISCARD		
□ RETURN (additional shipping charges may apply)		
Internal Use Only		
Approval:		
P.O.#:		
Date Received:		

Submission Information

Platform

Illumina Infinium MethylationEPIC BeadArrays for DNA derived from FFPE, frozen tissue, or fresh cells.

Sample Requirements:

Please ensure the DNA samples submitted meet the quality, concentration and volume requirements indicated below. **The BeadChip has a 8-sample per array format, please submit samples in sets of 8.**

- 1. Please provide at least 1µg of DNA derived from fresh cells or frozen tissue and at least 500ng of DNA derived from FFPE tissue. If your samples do not meet these minimum input requirements please contact Sandy Hsu (hsut@jwci.org) to discuss the options.
- 2. Sample concentration must be between $25 \text{ng}/\mu\text{L}$ and $100 \text{ng}/\mu\text{L}$. Please concentrate the sample if the DNA concentration is below $25 \text{ng}/\mu\text{L}$. DNA should be quantified by PicoGreen or similar fluorescent assay.
- 3. Sample must be suspended in nuclease-free water or 1X TE buffer; Volume ≥ 15 µl.
- 4. Removal of melanin contamination is highly recommended.
- 5. Samples should be contained in a 1.5 mL tube with the lid secured with parafilm. **Please clearly label each sample tube** with the following:
 - SampleID on the cap and side of each tube; SampleID should match the SampleID provided in the MethylationEPIC sample sheet.
 - Date and institution or PI initials on the side of each tube.
- 6. All samples will be quantified by PicoGreen upon arrival at JWCI. We will use $2\mu L$ of sample volume to determine DNA concentration.
- 7. An incomplete MethylationEPIC Sample Submission Form or MethylationEPIC sample sheet will result in a delay with your order.
- 8. The submitter is responsible for the samples submitted to the Sequencing Center including all necessary consent, HIPAA compliance, and any applicable specimen regulations.

Sample Shipping Instructions:

John Wayne Cancer Institute 2200 Santa Monica Blvd. Rm 128 Sequencing Center Santa Monica, CA 90404 Attn: Sequencing Center

- Please email shipment tracking information to Sandy Hsu (hsut@jwci.org)
- Samples should be shipped on enough dry ice to keep the sample frozen upon arrival at JWCI.

I have read and understand the above submission information and requirements

Signature:	
Date:	