



May 12, 2022

# GenomeTrakr WGS Protocol Collection and Workflow for MiSeq

Tina.Pfefer<sup>1</sup>, Julie Haendiges<sup>1</sup>, Maria Balkey<sup>1</sup>, Ruth Timme<sup>1</sup><sup>1</sup>US Food and Drug Administration

1

[dx.doi.org/10.17504/protocols.io.3b4l4bwyjvo5/v1](https://dx.doi.org/10.17504/protocols.io.3b4l4bwyjvo5/v1)**GenomeTrakr**Tech. support email: [genomeTrakr@fda.hhs.gov](mailto:genomeTrakr@fda.hhs.gov)

Tina.Pfefer

Here we have created a collection of all the protocols used for WGS using the MiSeq, in order, from sample extraction to NCBI submission.

## This collection has three sections:

- **Section 1: WGS Wet lab workflow for Illumina MiSeq**
- **Section 2: Dry lab workflow for sequence QC and NCBI submission - Direct Submission**
- **Section 3: Dry lab workflow for sequence QC and NCBI submission - PulseNet labs**

Associated protocols:

- [Querying the NCBI database for GenomeTrakr data \(protocols.io\)](#)
- [NCBI data curation protocol - SOP for editing GenomeTrakr submissions \(protocols.io\)](#)

DOI

[dx.doi.org/10.17504/protocols.io.3b4l4bwyjvo5/v1](https://dx.doi.org/10.17504/protocols.io.3b4l4bwyjvo5/v1)

Tina.Pfefer , Julie Haendiges, Maria Balkey, Ruth Timme 2022. GenomeTrakr WGS Protocol Collection and Workflow for MiSeq. **protocols.io**  
<https://dx.doi.org/10.17504/protocols.io.3b4l4bwyjvo5/v1>



GenomeTrakr, whole genome sequencing, enteric pathogens, surveillance, MiSeq

\_\_\_\_\_ protocol ,

Apr 05, 2022

May 12, 2022

60341

## Wet lab

### 1 WGS Wet lab workflow for Illumina MiSeq

### 2 DNA Extraction



Manual DNA Extraction using Qiagen DNeasy Blood and Tissue Kit  
by **Julie Haendiges**,  
US Food and Drug Administration

PREVIEW

RUN



### 3 DNA Quantification



DNA Quantification using the Qubit Fluorometer  
by **Julie Haendiges**,  
US Food and Drug Administration

PREVIEW

RUN



### 4 Library Preparation



Illumina DNA Prep (M) Tagmentation Library Preparation for use on an Illumina MiSeq Sequencer  
by **Julie Haendiges**,  
US Food and Drug Administration

PREVIEW

RUN



### 5 Sequencing



Procedure for Operation and Maintenance of the Illumina  
MiSeq for Whole Genome Sequencing  
**by Julie Haendiges,**  
**US Food and Drug Administration**

[PREVIEW](#)[RUN](#)

## Dry lab - Direct Submission

### 6 Dry lab workflow for sequence QC and NCBI submission - Direct Submission:

The following protocols are also included in a Springer Methods book chapter collection:  
[Utilizing the Public GenomeTrakr Database for Foodborne Pathogen Traceback \(protocols.io\)](#)

### 7 Check sequence quality:



Quality control assessment for microbial genomes:  
GalaxyTrakr MicroRunQC workflow  
**by Ruth Timme,**  
**US Food and Drug Administration**

[PREVIEW](#)[RUN](#)

### 8 Populate BioSample AND SRA metadata templates:



Guidance for populating GenomeTrakr metadata  
templates (BioSample and SRA)  
**by Ruth Timme,**  
**US Food and Drug Administration**

[PREVIEW](#)[RUN](#)

### 9 Submit sequence and metadata to NCBI:



NCBI submission protocol for microbial pathogen  
surveillance  
**by Ruth Timme,**  
**US Food and Drug Administration**

[PREVIEW](#)[RUN](#)

## 10 Update, retract, or replace these records in NCBI databases, if necessary:

[NCBI data curation protocol - SOP for editing GenomeTrakr submissions \(protocols.io\)](#)

Dry lab - PulseNet submission

## 11 Dry lab workflow for sequence QC and NCBI submission - PulseNet labs:

## 12 Check sequence quality:

Follow the CDC's guidance for assessing QC in BioNumerics

## 13 Populate NCBI template:



Populating NCBI template for submissions using  
BioNumerics v7.6  
by Maria Balkey

PREVIEW

RUN



## 14 Submit sequence and metadata to NCBI:

Follow the CDC's guidance for NCBI submissions through BioNumerics.

## 15 Update, retract, or replace these records in NCBI databases, if necessary:

[NCBI data curation protocol - SOP for editing GenomeTrakr submissions \(protocols.io\)](#)