

# CCR Genomic Data Sharing Plan

## Demographics

Investigator: \_\_\_\_\_

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

Date: \_\_\_\_\_  
Month/Day/Year

## Project Information

Project Name: \_\_\_\_\_

Planned enrollment/Sample Size: \_\_\_\_\_

### Collection Type

- Prospective
- Retrospective

## Data

### Source [check all that apply]

- Human
- Non-Human

### Sample Types [check all that apply]

- Germ line
- Somatic
- DNA
- RNA
- Microbiome
- Mitochondria
- Other, please specify : \_\_\_\_\_

**Type [check all that apply]**

***Array Data***

- Single Nucleotide Polymorphisms (SNP)
- Expression Array
- Methylation Array

***Genotypes***

- Array Derived Genotypes
- CNV Calls from Microarray
- CNV Calls Derived from Sequencing
- Somatic SNV(.MAF)
- Array CGH CNVs

***Sequencing***

- Whole Genome
- Whole Exome
- Targeted Genome
- Targeted Exome
- Whole Transcriptome
- Targeted Transcriptome
- Epigenomic Marks
- Sanger
- 16S rRNA

***Analysis***

- Association/Linkage
- Array Derived Expression
- RNA Seq Derived Expression
- Array Derived Methylation

***Other***

- Other, please specify : \_\_\_\_\_

**Data Form [check all that apply]**

- Individual
- Aggregate
- Not applicable

**Data Repository [check all that apply]**

- Array Express
- Cancer Genomics Hub
- ClinVar
- Database of Genotypes and Phenotypes (dbGaP)
- Database of Short Genetic Variants (dbSNP)
- Database of Genomic Structural Variation (dbVar)
- European Nucleotide Archive (ENA)
- GenBank
- Gene Expression Omnibus (GEO)
- Mouse Genome Informatics (MGI)
- Sequence Read Archive (SRA)
- Trace Archive
- WormBase
- Zebrafish Model Organism Database (ZFIN)
- Other, please specify : \_\_\_\_\_

**Companion Information [check all that apply]**

- Phenotype
- Exposure
- Clinical Data, please specify: \_\_\_\_\_
- Study/Protocol, please specify: \_\_\_\_\_
- Data collection instrument, please specify: \_\_\_\_\_
- Survey Tool
- Other, please specify : \_\_\_\_\_

## Submission Information

### Data Submission and Release Timeline [check all that apply]

- Following data cleaning, analysis, and quality checks
- At the time of publication
- Other, please specify : \_\_\_\_\_

### Data Use Limitations

- Not applicable, no human data
- No data use limitations
- Limited only by the terms of the Data Use Certification.
- Limited to health/medical/biomedical purposes, does not include the study of population origins or ancestry.
- Must be related to the specified disease.

### Approvals

	DATE	APPROVAL	NAME (Typed or Printed)	SIGNATURE
CCR GPA		<input type="checkbox"/> Yes <input type="checkbox"/> No		
CCR Scientific Director		<input type="checkbox"/> Yes <input type="checkbox"/> No*		

\* If No, please specify issues to be addressed: \_\_\_\_\_

Forms approved and signed off by the CCR Scientific Director should be returned electronically to the Investigator with a copy to the CCR GPA.