Bioinformatics for Disease

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Preface

This is a Quarto book.

To learn more about Quarto books visit https://quarto.org/docs/books.

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1 Molecular Sequencing Concepts and Technologies

Part I Data skills

In summary, this book has no content whatsoever.

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[1] 2

2 Common data visualization techniques

Part II

DNA

3 Aligning DNA sequences

4 DNA quality control

Here I'm listing some useful reference material.

Chapter 14

5 Variant calling

Part III

RNA

6 Aligning RNA sequences

7 RNA quality control

8 Differential expression analysis

9 Gene set enrichment analysis (GSEA)

Part IV Cancer Analyses

10 Cancer drivers

11 Driver enrichment

12 Driver enrichment

Part V Appendices

Here I'm listing some useful reference material.

 ${\rm Chapter}\ {\color{red} {14}}$

13 Reference versions

13.1 Downloading genome reference files

Species	Version	Command
human	GRCh38/h	$_{ m ig38wget}$
		https://hgdownload.soe.ucsc.edu/goldenPath/hg38/bigZips/hg38.fa.gz
human	GRCh37/h	m lg19wget
		https://hgdownload.soe.ucsc.edu/goldenPath/hg19/bigZips/hg19.fa.gz
mouse	GRCm39/r	mm%get
		https://hgdownload.soe.ucsc.edu/goldenPath/mm39/bigZips/mm39.fa.gz
mouse	GRCm38/r	mm liget
		https://hgdownload.soe.ucsc.edu/goldenPath/mm10/bigZips/mm10.fa.gz

13.2 Additional Resources

1. Broad Institute

14 Useful databases

14.1 Downloading genome reference files

Species	Version	Command
human	GRCh38/hg38	wget
	,	https://hgdownload.soe.ucsc.edu/goldenP
human	GRCh37/hg19	wget
		https://hgdownload.soe.ucsc.edu/goldenP
mouse	GRCm39/mm39	wget
	·	https://hgdownload.soe.ucsc.edu/goldenP
mouse	$\mathrm{GRCm}38/\mathrm{mm}10$	wget
	*	https://hgdownload.soe.ucsc.edu/goldenP

14.2 Additional Resources

1. Broad Institute

References