Check out the first project

Assignment invitation link:

https://classroom.github.com/a/84iMuKb5

\$ git clone git@github.com:coletrapnell-teaching/scRNAseq-starter-<your_gihub_id>.git

Repo structure

▼ 🛅 sc	i-RNA-seq-pipeline-scripts	۵	Today at 2:05 PM		Folder
7	assign-reads-to-genes.jun-pipeline-compatible.py	۵	Today at 1:10 PM	4 KB	Python Source
•	assign-reads-to-genes.jun-pipeline-compatible.sh	٥	Today at 1:10 PM	772 bytes	Shell Script
7	assign-reads-to-genes.py		Today at 1:10 PM	4 KB	Python Source
1	assign-reads-to-genes.sh	0	Today at 1:10 PM	764 bytes	Shell Script
**	count-rRNA-reads.sh	٥	Today at 1:58 PM	385 bytes	Shell Script
-	count-UMI-per-sample.sh	٥	Today at 1:10 PM	818 bytes	Shell Script
16	debug-assign-reads-to-genes.sh		Today at 1:10 PM	708 bytes	Shell Script
•	kallisto.sh	0	Today at 1:10 PM	339 bytes	Shell Script
ä	knee-plot-merge-samples.R	٥	Today at 1:10 PM	1 KB	Rez Source
Ä	knee-plot.R	٥	Today at 1:10 PM	1 KB	Rez Source
1	make-fasta-for-kallisto.sh	0	Today at 1:10 PM	618 bytes	Shell Script
	Nextera.v2.0.P7.oligo.rev.comp	0	Today at 1:10 PM	1 KB	Document
W	prelim-umi-count-stats.sh	٥	Today at 1:10 PM	555 bytes	Shell Script
	put-read1-info-in-read2.awk	٥	Today at 1:10 PM	2 KB	Document
	put-read1-info-in-read2.fancy-sample-mapping.awk		Today at 1:10 PM	4 KB	Document
1	put-read1-info-in-read2.fancy-sample-mapping.sh	۵	Today at 1:10 PM	678 bytes	Shell Script
W	put-read1-info-in-read2.sh	٥	Today at 1:10 PM	657 bytes	Shell Script
*	rmdup-and-make-split-bed.sh	٥	Today at 1:10 PM	532 bytes	Shell Script
	rmdup.awk	0	Today at 1:10 PM	1 KB	Document
1	run-bcl2fastq.sh	۵	Today at 1:10 PM	423 bytes	Shell Script
W	run-trim-galore-for-c-elegans.sh	٥	Today at 1:10 PM	370 bytes	Shell Script
*	run-trim-galore.sh	٥	Today at 1:10 PM	358 bytes	Shell Script
16	samtools-filter-sort.sh	0	Today at 1:10 PM	363 bytes	Shell Script
	sci-RNA-seq.P5.oligo.rev.comp	۵	Today at 1:10 PM	1 KB	Document
	sci-RNA-seq.RT.oligos	۵	Today at 1:10 PM	1 KB	Document
-	STAR-alignReads-8-cores.sh	٥	Today at 1:10 PM	610 bytes	Shell Script
16	STAR-alignReads-moreMem.sh	۵	Today at 1:10 PM	611 bytes	Shell Script
16	STAR-alignReads.sh	۵	Today at 1:13 PM	611 bytes	Shell Script
W	UMI-count-rollup.sh	۵	Today at 2:05 PM	289 bytes	Shell Script
🦸 so	ci-RNA-seq.pipeline.elegans.sh	٥	Today at 2:02 PM	13 KB	Shell Script