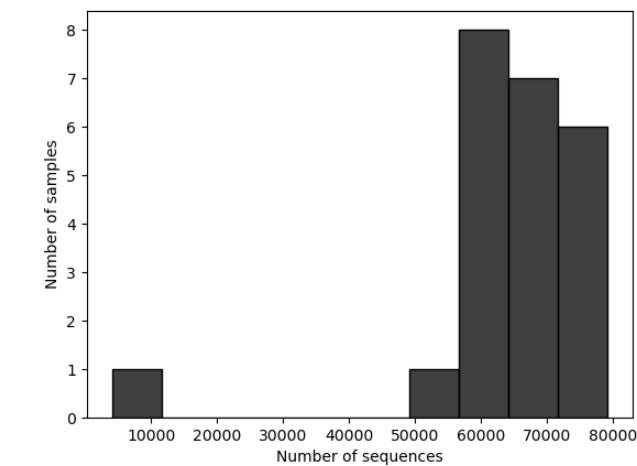


# Demultiplexed sequence counts summary

	forward reads	reverse reads
Minimum	4012	4012
Median	65675.0	65675.0
Mean	64081.434783	64081.434783
Maximum	79184	79184
Total	1473873	1473873

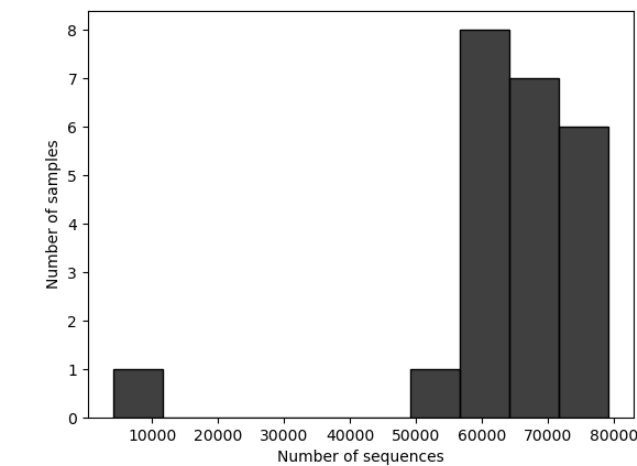
## Forward Reads Frequency Histogram



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(demultiplex-summary-forward.pdf)

## Reverse Reads Frequency Histogram



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(demultiplex-summary-reverse.pdf)

# Per-sample sequence counts

Total Samples: 23 (forward) 23 (reverse)

	forward sequence count	reverse sequence count
sample ID		

<b>EF1</b>	72278	72278
<b>EF2</b>	78211	78211
<b>EF3</b>	66507	66507
<b>EF4</b>	79184	79184
<b>EP1</b>	62950	62950
<b>EP2</b>	70408	70408
<b>EP3</b>	59581	59581
<b>EP4</b>	63589	63589
<b>EP5</b>	76552	76552
<b>KF1</b>	58106	58106
<b>KP1</b>	63420	63420
<b>KP2</b>	64301	64301
<b>KP3</b>	59816	59816
<b>KP4</b>	57701	57701
<b>KP5</b>	74346	74346
<b>KP6</b>	68343	68343
<b>NK</b>	4012	4012
<b>PK</b>	55046	55046
<b>TF1</b>	74620	74620
<b>TF2</b>	69194	69194
<b>TP1</b>	65675	65675
<b>TP2</b>	59767	59767
<b>TP3</b>	70266	70266

[Download as TSV \(per-sample-fastq-counts.tsv\)](#)