

Supplementary figure 2.501

```
$ ffq -l 2 GSM4769387 -o metadata.json
```

```
{
  "GSM4769387": {
    "accession": "GSM4769387",
    "supplementary_files": [
      {
        "filename": "GSM4769387_barcodes_S2.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM4769nnn/GSM4769387/suppl/GSM4769387_barcodes_S2.tsv.gz",
        "size": "15922"
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        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM4769nnn/GSM4769387/suppl/GSM4769387_features_S2.tsv.gz",
        "size": "303112"
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      {
        "filename": "GSM4769387_matrix_S2.mtx.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM4769nnn/GSM4769387/suppl/GSM4769387_matrix_S2.mtx.gz",
        "size": "43088259"
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    ],
    "platform": {
      "accession": "GPL24676",
      "title": "Illumina NovaSeq 6000 (Homo sapiens)"
    },
    "sample": {
      "SRS7325385": {
        "accession": "SRS7325385",
        "title": "O33_02_FL_S2",
        "organism": "Homo sapiens",
        "attributes": {
          "source_name": "human nHTBE cells",
          "biomaterial_provider": "Lonza CC-2540S",
          "tissue": "tracheal/bronchial scrapings",
          "cell_type": "Normal human tracheal bronchial epithelial",
          "isolate": "Batch 18TL134625, Tissue acquisition number 35825",
          "age": "65",
          "Sex": "male",
          "treatment": "untreated",
          "infection status": "SARS-CoV-2 infected, MOI 5, 24 hours post infection",
          "ENA-SPOT-COUNT": "232183638",
          "ENA-BASE-COUNT": "22986180162",
          "ENA-FIRST-PUBLIC": "2020-09-06",
          "ENA-LAST-UPDATE": "2020-11-04"
        },
        "experiment": "SRX9078405"
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  }
}
```