

Supplementary figure 2.500

```
$ ffq -l 2 GSM4769389 -o metadata.json
```

```
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    "accession": "GSM4769389",
    "supplementary_files": [
      {
        "filename": "GSM4769389_barcodes_S5.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM4769nnn/GSM4769389/suppl/GSM4769389_barcodes_S5.tsv.gz",
        "size": "12071"
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        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM4769nnn/GSM4769389/suppl/GSM4769389_features_S5.tsv.gz",
        "size": "303112"
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      {
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        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM4769nnn/GSM4769389/suppl/GSM4769389_matrix_S5.mtx.gz",
        "size": "33610735"
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    ],
    "platform": {
      "accession": "GPL24676",
      "title": "Illumina NovaSeq 6000 (Homo sapiens)"
    },
    "sample": {
      "SRS7325387": {
        "accession": "SRS7325387",
        "title": "O33_05_FL_S5",
        "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, 48 hours post infection",
          "ENA-SPOT-COUNT": "225964224",
          "ENA-BASE-COUNT": "22370458176",
          "ENA-FIRST-PUBLIC": "2020-09-06",
          "ENA-LAST-UPDATE": "2020-11-04"
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      "experiment": "SRX9078407"
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}
```