

Supplementary figure 2.388

```
$ ffq -l 2 GSM3738543 -o metadata.json
```

```
{
  "GSM3738543": {
    "accession": "GSM3738543",
    "supplementary_files": [
      {
        "filename": "GSM3738543_26y_F_PB_barcodees.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3738nnn/GSM3738543/suppl/GSM3738543_26y_F_PB_barcodees.tsv.gz",
        "size": "7201"
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        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3738nnn/GSM3738543/suppl/GSM3738543_26y_F_PB_genes.tsv.gz",
        "size": "257281"
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      {
        "filename": "GSM3738543_26y_F_PB_matrix.mtx.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3738nnn/GSM3738543/suppl/GSM3738543_26y_F_PB_matrix.mtx.gz",
        "size": "6719394"
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    ],
    "platform": {
      "accession": "GPL18573",
      "title": "Illumina NextSeq 500 (Homo sapiens)"
    },
    "sample": {
      "SRS4698196": {
        "accession": "SRS4698196",
        "title": "26y_F_PB",
        "organism": "Homo sapiens",
        "attributes": {
          "source_name": "Blood",
          "tissue": "Blood",
          "cell type": "NK cells",
          "sorting population": "Lin-CD7+",
          "genotype/variation": "healthy donor",
          "gender": "female",
          "ENA-SPOT-COUNT": "175269685",
          "ENA-BASE-COUNT": "17176429130",
          "ENA-FIRST-PUBLIC": "2019-07-16",
          "ENA-LAST-UPDATE": "2019-07-16"
        },
        "experiment": "SRX5763727"
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    }
  }
}
```