

## Supplementary figure 2.384

```
$ ffq -l 2 GSM3738539 -o metadata.json
```

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      {
        "filename": "GSM3738539_26y_F_BM_barcodes.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3738nnn/GSM3738539/suppl/GSM3738539_26y_F_BM_barcodes.tsv.gz",
        "size": "6433"
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        "size": "5953131"
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    "platform": {
      "accession": "GPL18573",
      "title": "Illumina NextSeq 500 (Homo sapiens)"
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    "sample": {
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        "title": "26y_F_BM",
        "organism": "Homo sapiens",
        "attributes": {
          "source_name": "Bone marrow",
          "tissue": "Bone marrow",
          "cell type": "NK cells",
          "sorting population": "Lin-CD7+",
          "genotype/variation": "healthy donor",
          "gender": "female",
          "ENA-SPOT-COUNT": "189789087",
          "ENA-BASE-COUNT": "18599330526",
          "ENA-FIRST-PUBLIC": "2019-07-16",
          "ENA-LAST-UPDATE": "2019-07-16"
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      "experiment": "SRX5763723"
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}
```