

## Supplementary figure 2.394

```
$ ffq -l 2 GSM3738540 -o metadata.json
```

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{
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      {
        "filename": "GSM3738540_25y_M_BM_barcodes.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3738nnn/GSM3738540/suppl/GSM3738540_25y_M_BM_barcodes.tsv.gz",
        "size": "4716"
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        "size": "3196828"
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    ],
    "platform": {
      "accession": "GPL18573",
      "title": "Illumina NextSeq 500 (Homo sapiens)"
    },
    "sample": {
      "SRS4698193": {
        "accession": "SRS4698193",
        "title": "25y_M_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": "male",
          "ENA-SPOT-COUNT": "118598973",
          "ENA-BASE-COUNT": "11622699354",
          "ENA-FIRST-PUBLIC": "2019-07-16",
          "ENA-LAST-UPDATE": "2019-07-16"
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      "experiment": "SRX5763724"
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}
```