

Supplementary figure 2.390

```
$ ffq -l 2 GSM3739218 -o metadata.json
```

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        "filename": "GSM3739218_ET05.barcodes.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3739nnn/GSM3739218/suppl/GSM3739218_ET05.barcodes.tsv.gz",
        "size": "21283"
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        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3739nnn/GSM3739218/suppl/GSM3739218_ET05.matrix.mtx.gz",
        "size": "33239165"
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    ],
    "platform": {
      "accession": "GPL16791",
      "title": "Illumina HiSeq 2500 (Homo sapiens)"
    },
    "sample": {
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        "accession": "SRS4701336",
        "title": "10x.ET05",
        "organism": "Homo sapiens",
        "attributes": {
          "source_name": "ET05, scRNA-seq",
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          "targeted_mutation": "CALR exon9 A382Rfs*58 (c.1144_1215delins67)",
          "Sex": "male",
          "age": "17",
          "diagnosis": "ET",
          "pb blast count": "0%",
          "bm blast count": "1%",
          "bm fibrosis": "MF-0/3",
          "treatment": "None",
          "ENA-SPOT-COUNT": "148890936",
          "ENA-BASE-COUNT": "14591311728",
          "ENA-FIRST-PUBLIC": "2019-06-02",
          "ENA-LAST-UPDATE": "2019-06-02"
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        "experiment": "SRX5767042"
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}
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