

## Supplementary figure 2.209

```
$ ffq -l 2 GSM3378059 -o metadata.json
```

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{
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      {
        "filename": "GSM3378059_ZFN_barcodes.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3378nnn/GSM3378059/suppl/GSM3378059_ZFN_barcodes.tsv.gz",
        "size": "16037"
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        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3378nnn/GSM3378059/suppl/GSM3378059_ZFN_matrix.mtx.gz",
        "size": "31416853"
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    ],
    "platform": {
      "accession": "GPL18573",
      "title": "Illumina NextSeq 500 (Homo sapiens)"
    },
    "sample": {
      "SRS3747198": {
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        "title": "ZFN",
        "organism": "Homo sapiens",
        "attributes": {
          "source_name": "FACS-sorted primitive and progenitors HSPC in ratio 1 to 1"
        },
        "time point": "24h upon nuclease delivery",
        "cell type": "Cord blood derived CD34+ cells",
        "ENA-SPOT-COUNT": "159684897",
        "ENA-BASE-COUNT": "15649119906",
        "ENA-FIRST-PUBLIC": "2019-03-02",
        "ENA-LAST-UPDATE": "2019-03-02"
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      "experiment": "SRX4649197"
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}
```