

Supplementary figure 2.215

```
$ ffq -l 2 GSM3378056 -o metadata.json
```

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{
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    "supplementary_files": [
      {
        "filename": "GSM3378056_RNPhs_barcodes.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3378nnn/GSM3378056/suppl/GSM3378056_RNPhs_barcodes.tsv.gz",
        "size": "9504"
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        "filename": "GSM3378056_RNPhs_genes.tsv.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3378nnn/GSM3378056/suppl/GSM3378056_RNPhs_genes.tsv.gz",
        "size": "284858"
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      {
        "filename": "GSM3378056_RNPhs_matrix.mtx.gz",
        "url": "ftp.ncbi.nlm.nih.gov/geo/samples/GSM3378nnn/GSM3378056/suppl/GSM3378056_RNPhs_matrix.mtx.gz",
        "size": "14590087"
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    ],
    "platform": {
      "accession": "GPL18573",
      "title": "Illumina NextSeq 500 (Homo sapiens)"
    },
    "sample": {
      "SRS3747195": {
        "accession": "SRS3747195",
        "title": "HS RNP",
        "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": "144813244",
        "ENA-BASE-COUNT": "14191697912",
        "ENA-FIRST-PUBLIC": "2019-03-02",
        "ENA-LAST-UPDATE": "2019-03-02"
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      "experiment": "SRX4649194"
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}
```