## **STA323**

# Big Data Analysis Software and Application (Hadoop or Spark) Report on Assignment 2

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Question 1. The protein sequence data is in the remote server /shareddata/data/as2/Q1\_data. Here, fasta is a widely used file format in bioinformatics. In the fasta file, the line starting with '>' is the header of the sequence, i.e. comment for the protein sequence.

在题目的最后声明忽略氨基酸序列的拼接问题,但在此之前我已经实现了氨基酸序列的拼接,故实现此部分的注释和代码以加删除线的形式呈现。

(1). Calculate the frequency of each type of amino acid with Spark Dataframe API. [1 point] Answer:

首先筛选出注释行(以'>'开头的行),用 is\_header 列标记之<del>,并用 is\_header 的前缀和 group 标记氨基酸的分组(即其属于第几个蛋白质)</del>。用 line\_num 标记行号,然后删去注释 行。

```
fasta_df = fasta_df.withColumn("is_header", when(col("value").startswith(">"), 1).otherwise(0))
fasta_df = fasta_df.withColumn("line_num", monotonically_increasing_id())
fasta_df = fasta_df.withColumn("group", expr("sum(int(is_header)) over (order by line_num)"))
filtered_df = fasta_df.filter(col("is_header") == 0)
```

value is_	neader lin		
MEEITQIKKRLSQTVRL	0	1	1
KKEVVAVAKKEEVLKKE	0	2	1
DIVPQMRDVSLPPKEEE	0	3	1
SLPPKKDEEIVCEKKEV	0	4	1
KEHEEKETFVVLKKEII	0	5	1

可以用下列代码将同组的氨基酸序列拼接。

merged\_df = filtered\_df.groupBy("group").agg(concat\_ws("", collect\_list("value")).alias("sequence"))

使用 explode 函数展开各行并统计,即得各氨基酸出现的频数。 amino\_acid\_df = merged\_df.select(explode(split(col("value"), "")).alias("amino\_acid")) frequency\_df = amino\_acid\_df.groupBy("amino\_acid").count().orderBy("amino\_acid")

amino a	cidl	count
+	+	+
1	AL	3223081
i	В	6
İ	cl	564455
İ	D :	2216904
1	E	2674664
Ì	F	985877
İ	G :	2653426
Ì	H	628384
1	II:	1726915
İ	K   :	1684031
Ì	LI:	2851645
1	M	467474
1	NI:	1316889
Ì	PI:	2097950
1	QI:	1422769
1	RI:	1789613
1	SI:	2747798
1	TI:	2795042
1	VI:	2760761
1	W	351166
1	X	1347
1	Y	823096
1	ZI	2

(2). Complete the above work with Spark RDD API. [1 point]

### Answer:

```
使用 RDD 实现如前所述的流程即可。
def is header(line):
    return line.startswith(">")
def assign_group(iterator):
    group = 0
    for line in iterator:
        if is header(line):
           group += 1
        yield (group, line)
grouped_rdd = fasta_rdd.mapPartitions(assign_group)
filtered_rdd = grouped_rdd.filter(lambda x: not is_header(x[1]))
merged_rdd = filtered_rdd.groupByKey().mapValues(lambda lines: "".join(lines))
amino_acid_rdd = merged_rdd.flatMap(lambda x: list(x[1]))
frequency_rdd = amino_acid_rdd.map(lambda x: (x, 1)).reduceByKey(lambda a, b: a + b).sortByKey()
                                        Frequency of each amino acid:
                                        A 3223081
                                        B 6
                                        C 564455
                                        D 2216904
                                        E 2674664
                                        F 985877
                                        G 2653426
                                        H 628384
                                        I 1726915
                                        K 1684031
                                        L 2851645
                                        M 467474
                                        N 1316889
                                        P 2097950
                                        Q 1422769
                                        R 1789613
                                        5 2747798
                                        T 2795042
                                        V 2760761
                                        W 351166
                                        X 1347
                                        Y 823096
                                        Z 2
```

(3). Count the number of a specific sequence motif "STAT" with Spark. [2 points]

#### Answer

统计每行出现的 STAT 字样的次数并求和即可,得到共 2052 个。(如果考虑拼接则数量稍多一些)

```
 stat\_count\_df = merged\_df.withColumn("stat\_occurrences", (size(split(col("value"), "STAT")) - 1)) \\ stat\_count\_df.show(5) \\ total\_stat\_count = stat\_count\_df.agg(\{"stat\_occurrences": "sum"\}).collect()[0][0] \\
```

Question 2. Here is a dataset related to the online open courses. The course.csv contains the information about the courses while instructors.csv contains the information about the instructors.

(1). Join the 2 dataframes by instructors\_id, use the inner join function. [2 points]

#### Answer

```
instructor_df_renamed = instructor_df.withColumnRenamed("id",
"instructor_id").withColumnRenamed("title", "instructor_title")
joined_df = course_df.join(
    instructor_df_renamed,
    instructor_df_renamed.instructor_id == course_df.instructors_id, how = "inner"
)
joined_df.drop(joined_df["instructors_id"])
注意处理重名的列即可。
```

(2). Use SQL (in pyspark SQL) to show the display\_name and job\_title of the instructor, who has the highest course rating among all courses that are related to "spark". [2 points]

#### Answer:

```
首先创建视图。
course_df.createOrReplaceTempView("course")
instructor_df.createOrReplaceTempView("instructor")
joined_df.createOrReplaceTempView("course_with_instructor")
     依照题意筛选排序即可。
q2_2_df = spark.sql("""
   select
       display name,
       job_title,
       rating
      course_with_instructor
   where
       title like "%spark%"
       created >= "2018-01-01 00:00:00"
   and
       rating = (
          select
             max(rating)
          from
              course_with_instructor
          where
              title like "%spark%"
              created >= "2018-01-01 00:00:00"
   order by
       rating desc
q2_2_df.show(truncate = False)
                      |display_name|job_title
                      |Deby Coles |Sewer, Artist, Crafter and Instructor | 4.6432705 |
```

(3). Use SQL (in pyspark SQL) to select all courses that are contains "interview" or "interviews" and then sorted by course\_rating in descending order and created in descending order (newest first). In this task, the course rating should be firstly rounded to one decimal place. [2 points]

## Answer:

```
依照题意,注意将 title 小写化进行匹配,编写 SQL 即可。这里仅选中了必要的列。
q2_3_df = spark.sql("""
    select
        id,
        title,
        created,
        round(rating, 1) AS rating
    from
        course
    where
        lower(title) like "%interview%"
    order by
        rating desc,
        created desc
""")
q2_3_df.show(truncate = False)
```

id	title	created	rating
48869	926 Interview Oriented Data Structure Arrays & Linked List C C-	++ 2022-09-17T17:57:14Z	5.0
43094	400 CATIA V5 FOR JOBS INTERVIEW	2021-09-20T12:54:23Z	5.0
48291	150 Réaliser des interviews au rendu professionnel (PARTIE 2)	2022-08-12T14:54:06Z	4.9
47228	394 "The ""BigTech"" System Design Interview Bootcamp"	2022-06-07T14:53:40Z	4.9
44994	176 Power BI Interview Questions and Answers	2022-01-17T11:08:03Z	4.9