lecture 7: Dask

wbg231

January 2023

1 review of the story so far

- started at file systems
 - they are unstructured collections of files
 - there are many custom ways to store data
 - very flexible
 - there is no built in parallelism
- relational data bases
 - restrict the structure of data to relations
 - has a standard interface (SQL)
 - somewhat flexible
 - not easy to parallelize
- map reduce and HDFS
 - data is less structured than RDBMS
 - restrict coding to map and reduce functions
 - very parallel
- spark
 - structured data like RDBMS
 - distributed storage (HDFS)
 - standard-ish interface (sql and spark-API)
 - very parallel

spark is good

- spark integrates well with java based tools like hadoop
- spark is grate at working with data frames and SQL like data processing
- it is 10 years old meaning the implementation is mature and stable
- after RDBMS and SQl it is likely the most used software for data analysis

what is spark not good at

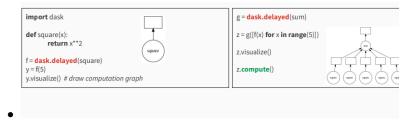
- data that does not fit nearly into an RDD or data frame model
- it is not super well suited for working with the python SciPy stack
- modern machine learning is all done using SciPY
- spark is only cluster based what if the data you are working with is to big to hold in RAM but don't need a cluster?

Dask

- Dask is python based distributed computation
- has a lot in common with spark
- has computation graphs similar to sparks linage graphs
- has delayed computation much like RDD and data frames in spark
- collection based interface also has spark like data frames
- some key differences between spark and dask are that
 - 1. dask prioritizes array based Numpy like computation
 - 2. it is designed to support single machine out of core (ie parallel or multi thread) use

delayed computation and task graphs

• dask builds complex computation by composing deferred computation into task graphs



- the left shows how a delayed computation square function can be defined
- the right shows how we can pass that computation to a delayed sum function and look at it in parallel

collections in dask

- bags
 - distributed collections of arbitrary structured data. similar to an RDD in spark
- data frames distributed collection of structured tabular data, most similar to a spark data frame (but built on pandas instead of using RDDs)
- arrays: n dimensional distributed numpy arrays.

collection interface bags

- bags are broadly similar to spark RDDs. think of them kinda like a list in python
- they are un ordered collections of generic python objects. portioned into subsets
- they implement basic operations like map filter join sum etc
- a good choice for initial preprocessing and structured objects
- if your data is tabular or array based this is likely not a good data structure

dask bags vs spark RDDs

- both partition a collection of objects across multiple machines
- both are immutable
- all elements of an rdd must be of the same type
- bags are untyped so the contents can be mixed types (but this should be avoided as it slows down processing)
- \bullet a common dask work flow is raw data \to bags \to data frames \to deeper analysis
- the earlier you reduce the size of your data the better, as that is less data moving through the system
- we prefer maps and filters on bags over data frame manipulations when simplifying data
- but in general bag operations are slower than those on data frames, since because bags have so few restrictions it is hard to optimize code for them

constructing collections

- there are many ways to construct a collection in dask
- db.from_sequence.([logad(f) for f in files]). all files are loaded first in vanilla python lists and then put in a bag
- db.from_sequence(files).map(load) file names are distributed into a bag, the load function is then applied to each in parallel
- so the second is faster
- in general try to load data using collections when possible

bag folding vs grouping

- try to avoid using group by on bags, this requires inter worker communications which is slow. there are wide dependencies with this method
- use fold or fold by if possible
- this method has similar benefits to a combiner in map reduce, that is it does local aggregation fit to reduce data shuffling
- fold by required a key function and binary operation
- a key function maps elements to a key (think of this as the group)
- a binary operation reduces within the group (think of this as group aggregation)
- a binop and reducer in map reduce are not the same however
- bag binop only sees two values at a time not a list like in map reduce
- output of a binop must match the input type unlike in map reduce
- $\bullet\,$ this can become tricky if your bag elements are structured

collection interface data frames

- they are similar to spark dataframes (they use pandas internally through)
- parallelism ie partitioning is over rows in dask data frames not cols
- these are a good choice for data that can naturally be split into different files. like multiple log records
- managing partitions in dask is important.

- your data may change through the computation graph if this is the case, you may end up with nearly empty partitions
- try to keep your partitions are full and balanced as possible
- dask makes you work harder than spark to get it working well

collections interface arrays

- dask arrays work like those in numpy
- parallelism is not limited to rows, can define chunks alone each dimension
- large arrays assembled implicitly from smaller arrays
- most numpy operations work automatically

where chunks fail

- not good for sorting between chunks (but can sometimes just take the top k elements from each chunk and use those)
- operations where the output size changes like masking operations are hard
- linear algebra operations can be tough

how spark is commonly used

- have 10 models for audio segmentation to compare
- have 2000 audio recordings in a dataset, that is 20,000 model outputs
- the model outputs are a sequence of time intervals
- Model evaluator compares reference annotation to estimate annotation for one track, and produces a dictionary of scores along different metrics.
 Takes a few seconds to run for each track.
- What I want: a DataFrame containing: model id, recording id, [scores for each metric]

solution

- store model output as separate files on disk
- creat a delayed function to map filenames to scores (calls the evaluator)
- crate a bag from the delayed function
- convert the bag to a data frame and save it

why is this better

- all computation is done using basic python functions (ewe don need to re-write our functions for dask)
- the problem it's self is parallel, so there is a lot of mapping but little reducing that needs to be done
- we like this over spark because we dont have to change any of our data structures or code to fit it

working with large numerical data

- csv and parquet files are not a good option here. (like a single really big matrix)
- collections of files npy or npz files can work well
- Hierarchical data format (HD5) could work well as well

HD5

• basically a new file system within a file (Hierarchical) Directory structures

does dask replace spark

- it kind of depends on use case
- pros for dask
 - works well with SciPY
 - works well with dense multi dimensional data
 - works well with custom algorithms and gpus
- pros for spark
 - more stable
 - more high level, you do not need to think as much about computation graphs or partitions
 - faster for data frame analysis
 - better support for large graph data

HPC

• hpc greene is less restrictive than dataproc

•