

A Pipeline for Processing Images in iDigBio to Train Convolutional Neural Networks

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iDigBio data and portal

Making data and images of millions of biological specimens available on the web

115,879,282

Specimen Records

27,335,463

Media Records

1,585

Recordsets

Search the Portal



Why digitization matters

More about what we do and why



Digitization

Learn, share and develop best practices



Sharing Collections

Documentation on data ingestion



Working Groups

Join in, contribute, be part of the community

iDigBio Home Portal Home Search Records Tutorial Our Data Research Tools Feedback

Search Records [Help](#) [Reset](#)

search all fields

Must have image Must have map point

Filters Mapping Sorting Download

Add a field Clear

Kingdom: Present Missing

Scientific Name: Present Missing [Add EOL Synonyms](#)

Date Collected: Start: End: Present Missing

Scroll To Bottom

[List](#) [Labels](#) [Images](#) [Recordsets](#)

Total: 20,495,711

Family	Scientific Name	Date Collected	Country	Institution Code	Basis of Record	Columns
no data	"Ambocoelia" sp.	no data	United States	MCZ	PreservedSpecimen	view
no data	"Ambocoelia" sp.	no data	United States	MCZ	PreservedSpecimen	view
Apterontidae	"Apterontus" apurensis	1981-01-08	Brasil	Instituto Nacional de Pesquisas da...	PreservedSpecimen	view
Apterontidae	"Apterontus" apurensis	1986-02-28	Brasil	Instituto Nacional de Pesquisas da...	PreservedSpecimen	view

iDigBio storage back-end

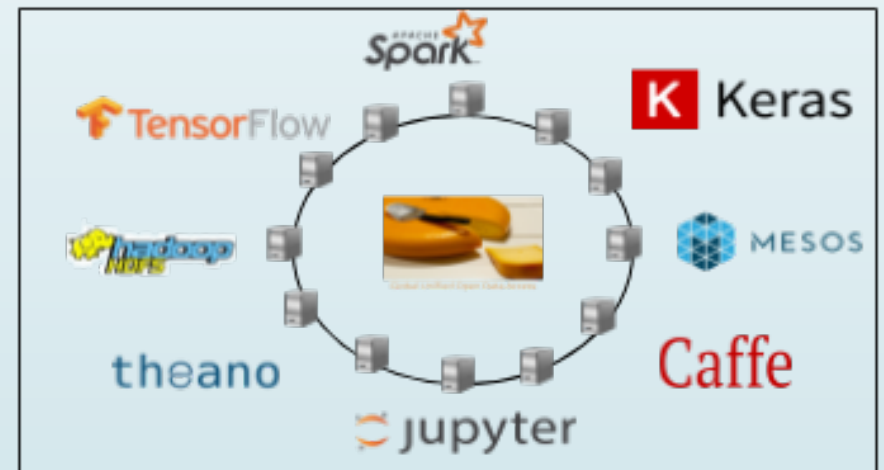
- ▶ 115+ million specimen records
- ▶ 27+ million media records
- ▶ 120 terabytes of media files
- ▶ Hosted on Ceph Distributed storage system
- ▶ Opportunity: machine learning processing near data



iDigBio Database of Images

GUODA – Global Unified Open Data Access

- ▶ Apache Mesos – distributed systems kernel
- ▶ Apache Spark – fast and general engine for large-scale data processing
- ▶ Jupyter Notebook Interface for Python and R
- ▶ Installed frameworks for deep learning –
 - ▶ TensorFlow
 - ▶ Keras
 - ▶ Theano
 - ▶ Caffe



GUODA cluster

A dark grey arrow points to the right from the left edge of the slide. Several thin, light blue lines curve downwards from the arrow's tip towards the bottom left corner of the slide.

Applications of Artificial Intelligence (AI)

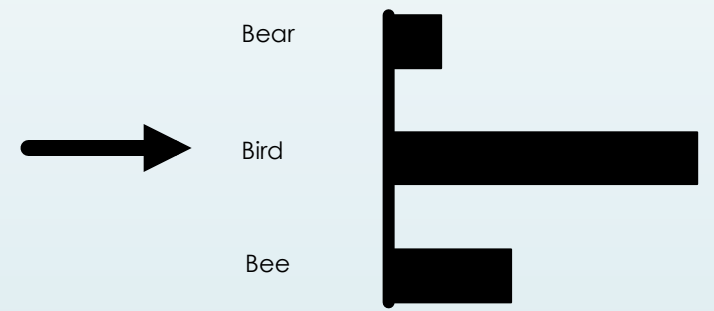
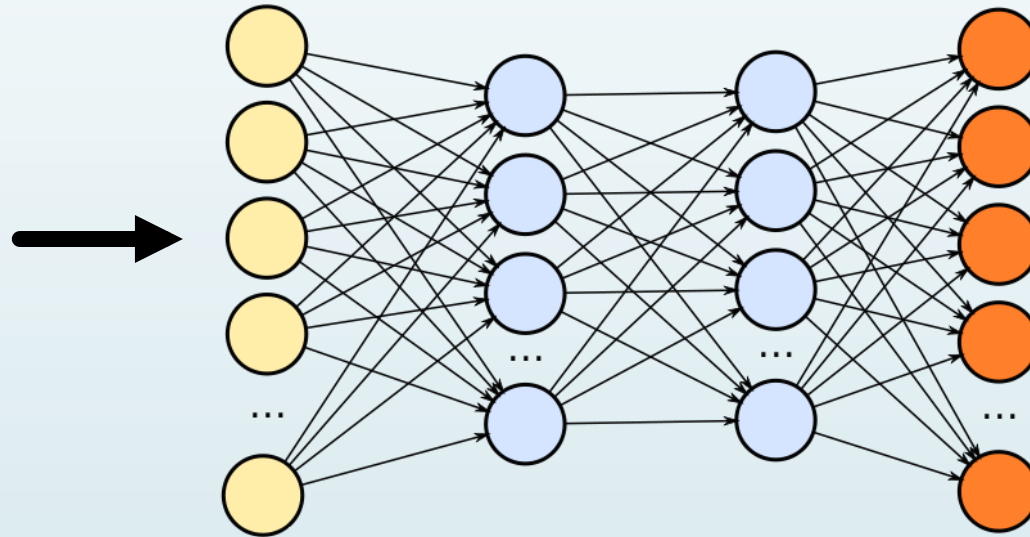
- ▶ Decision-making
- ▶ System Identification and Control
 - ▶ Vehicle control, process control, natural resource management
- ▶ Pattern Recognition
 - ▶ Face identification, object recognition, etc.
- ▶ Classification
 - ▶ Image classification, e-mail spam filtering
- ▶ **Emerging applications of AI for specimen images?**

Use Case – Find Mercury Contamination on Herbarium Specimens

- ▶ Mercury salt used to specimens as insecticides
- ▶ Mercury marking is visible on the specimens
- ▶ Mercury vapor and mercury compounds pose threat to human health.
- ▶ Mercury contaminated images can be classified using convolutional neural networks.
- ▶ E. Schuettpelz, P. Frandsen, R. Dikow, A. Brown, S. Orli, M. Peters, A. Metallo, V. Funk, L. Dorr. (2017). Applications of deep convolutional neural networks to digitized natural history collections. *Biodiversity Data Journal*. 5. e21139. 10.3897/BDJ.5.e21139.



Convolutional Neural Networks (CNN)



Accessing iDigBio Images

Search Records

search all fields

Must have media Must have map point

Filters Mapping Sorting Download

Add a field Clear

Scientific Name: [Add EOL Synonyms](#)

Present Missing

Date Collected: Start: End:

Present Missing

Country:

Present Missing

Total: 105,768,146

Family	Scientific Name	Date Collected	Country	Institution Code	Basis of Record	Columns
Hamamelidaceae	"Acer" (Liquidambar) lesqueureuxi	no data	United States	UCMP	FossilSpecimen	view
Hamamelidaceae	"Acer" (Liquidambar) lesqueureuxi	no data	United States	UCMP	FossilSpecimen	view
no data	"Ameea" pseudocrispula (Temb...	1976/01/01	Deutschland	no data	FossilSpecimen	view
no data	"Ameea" pseudocrispula (Temb...	1976/01/01	Deutschland	no data	FossilSpecimen	view

```
In [1]: from urllib.request import urlopen
from io import BytesIO
from PIL import Image
from imagehash import phash
from pyspark.sql import functions as F
from pyspark.sql import types as T
from pyspark.ml.linalg import Vectors

import numpy as np

In [2]: df = sqlContext.read.parquet("/guoda/data/idigbio-media-20171112T013207.parquet")

In [9]: df2 = df.sample(False,0.005,45) # to select small dataframe for faster execution
df.count()
df.printSchema()

root
 |-- accessuri: string (nullable = true)
 |-- data: struct (nullable = true)
 |   |-- Iptc4xmpExt:CountryCode: string (nullable = true)
 |   |-- Iptc4xmpExt:CountryName: string (nullable = true)
 |   |-- Iptc4xmpExt:ProvinceState: string (nullable = true)
 |   |-- Iptc4xmpExt:WorldRegion: string (nullable = true)
 |   |-- ac:accessURI: string (nullable = true)
 |   |-- ac:associatedSpecimenReference: string (nullable = true)
 |   |-- ac:attributionLogoURL: string (nullable = true)
 |   |-- ac:bestQualityAccessURI: string (nullable = true)
 |   |-- ac:bestQualityFormat: string (nullable = true)
 |   |-- ac:caption: string (nullable = true)
 |   |-- ac:captureDevice: string (nullable = true)
 |   |-- ac:comments: string (nullable = true)
 |   |-- ac:digitizationDate: string (nullable = true)
 |   |-- ac:furtherInformationURL: string (nullable = true)
 |   |-- ac:goodQualityAccessURI: string (nullable = true)
 |   |-- ac:hashFunction: string (nullable = true)

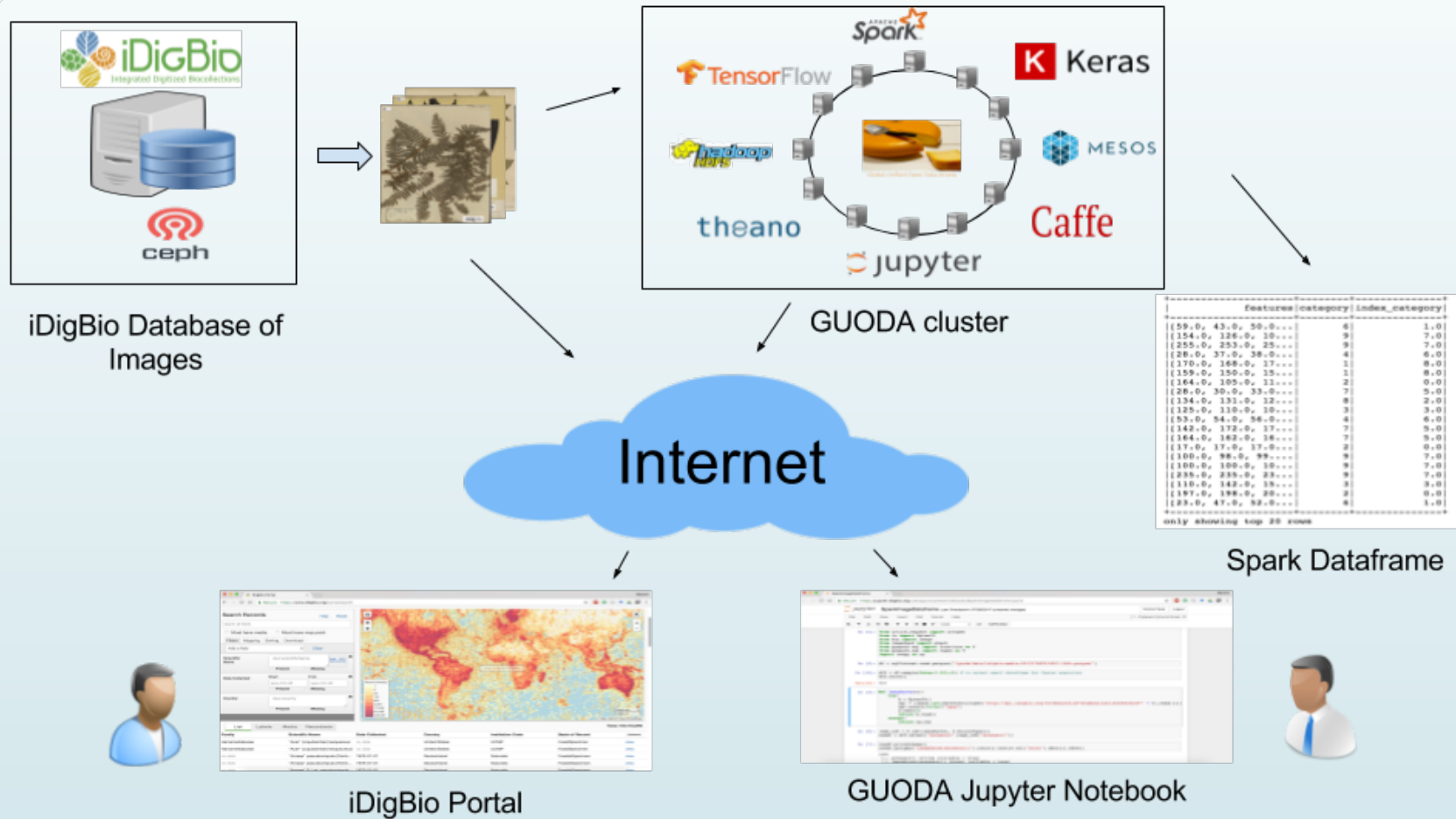
In [40]: data_df.first()[1]
```

Spark dataframe of iDigBio media

```
+-----+-----+-----+-----+
|          accessuri |contaminated|image_nparray(accessuri) |          vector_images|
+-----+-----+-----+-----+
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           1|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
|http://collection...|           0|[255.0, 216.0, 25...|[255.0,216.0,255....|
+-----+-----+-----+-----+
```

only showing top 20 rows

Image Processing Pipeline



Opportunities for collaboration

Founded in 2002

Focused on
researchers and
institutions on the
Pacific Rim

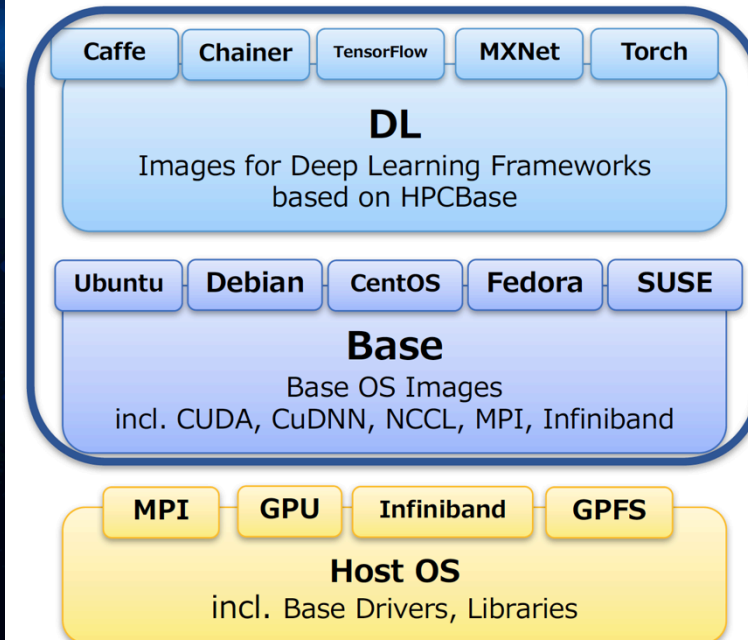
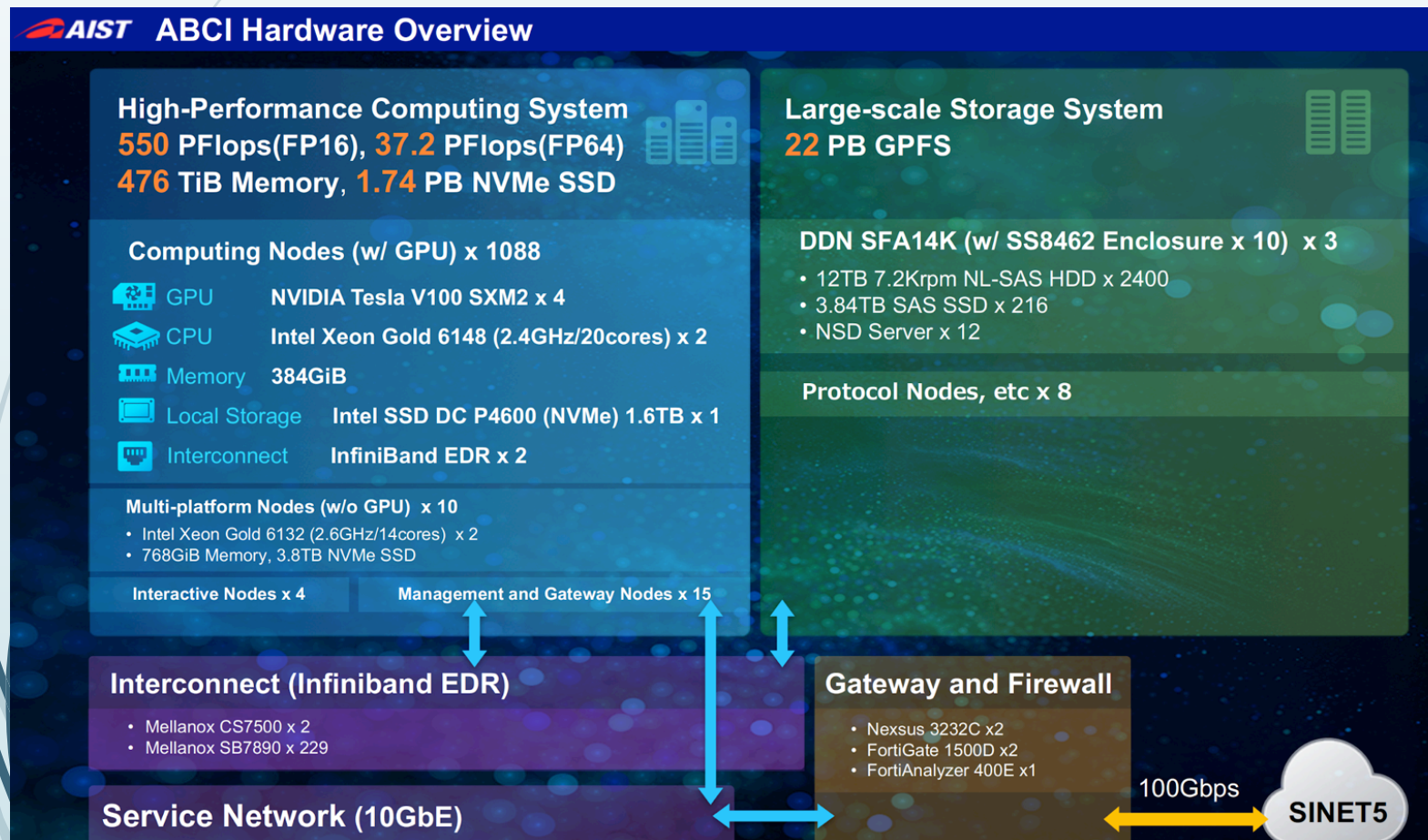
Open Community of
Practice

Engages “Long Tail”
science communities



AI-focused high-performance clusters

ABCI – top #5 in the world; similar system at NCHC, Taiwan





Future Work



- ▶ Join specimen metadata from iDigBio with images in dataframes which can serve as image and label pairs to train more exciting deep learning models
- ▶ Build another audio signal processing pipeline for existing audio data in iDigBio
- ▶ Make this pipeline portable and make it compatible to run on infrastructures such as NSF's XSEDE, AIST's ABCI

Acknowledgements

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