



NoSQL approaches in GnplS



Elasticsearch @ URGI Feedback



Data Discovery

- Links to datasets, by metadata
- Loose integration (keywords)
- Data model
 - Simple
 - Easy to implement and feed
- Ex:
 - GnpIS Google-like search
 - transPLANT search
 - WheatIS
 - IFB-Elixir
- Data set building, Data Mining
 - Combine data from different sources
 - Strong integration
 - Data model
 - Rich
 - Complex to model, implement and feed
 - Ex: GnpIS.Ephesis



Data discovery system

- GnplS, transPLANT, WheatIS searches
- Google like, full text, filters
- <http://www.transplantdb.eu>

CURRENT SEARCH
Found 98218 results

FILTER BY DATABASE:
• GnplS (98218)

FILTER BY DATA TYPE:
• SEQUENCE FEATURE (97996)
• ACCESSION (221)
• EXPERIMENT (1)

FILTER BY SPECIES:
• Vitis vinifera (97996)
• Vitis vinifera subsp vinifera cv.
Riesling (194)
• Vitis vinifera subsp vinifera cv.
Riesling italico (6)
• Vitis vinifera subsp vinifera cv.
Riesling bleu (Collection
Oberlin) (3)

Database entries Resources transPLANT web

Search » Search results » *vitis riesling*

Search

Search results

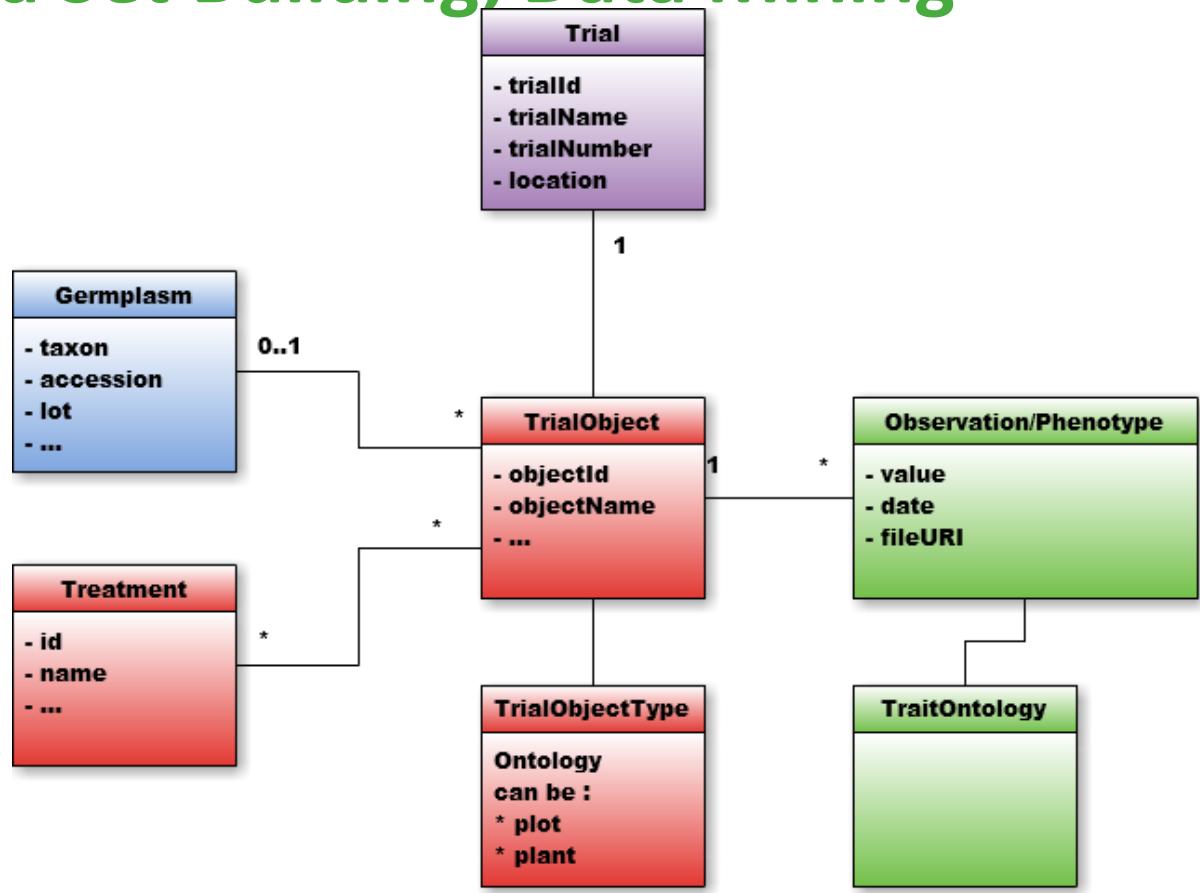
ACCESSION_31303
ACCESSION ... GnplS ... ACCESSION_31303 ... Riesling B2044 is a *Vitis vinifera* subsp *vinifera* cv. Riesling accession (status: Maintained) maintained by GRAPEVINE's GRC (managed by 'INRA Vassal Montpellier'), held by 'INRA UMR-SVQV Colmar' ... *Vitis vinifera* subsp *vinifera* cv. Riesling ... UMR Santé de la Vigne et Qualité du Vin, INRA-Colmar

ACCESSION_31346
ACCESSION ... GnplS ... ACCESSION_31346 ... Riesling B2185 is a *Vitis vinifera* subsp *vinifera* cv. Riesling accession (status: Maintained) maintained by GRAPEVINE's GRC (managed by 'INRA Vassal Montpellier'), held by 'INRA UMR-SVQV Colmar' ... *Vitis vinifera* subsp *vinifera* cv. Riesling ... UMR Santé de la Vigne et Qualité du Vin, INRA-Colmar

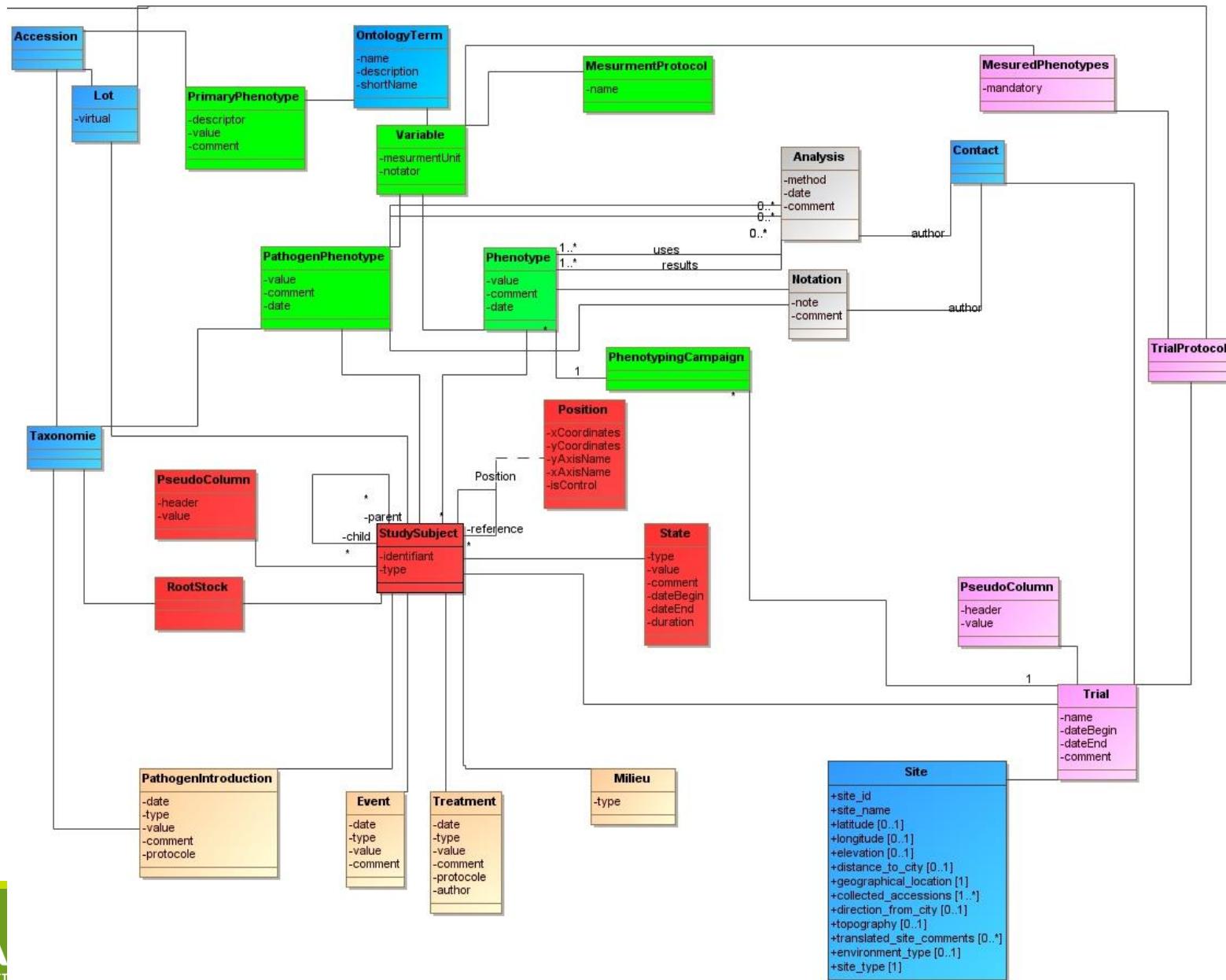
ACCESSION_31388

Data Set Building, Data Mining

- Phenotyping data
- Conceptual model

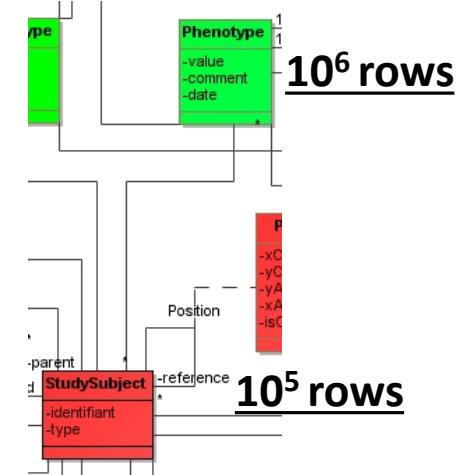


GnplIS.Ephesis Implementation



GnplS.Ephesis V1 Performances

- StudySubject **join** Phenotype
 - Too slow, too big
- Solution
 - Denormalisation, aggregation



10⁶ rows

10⁵ rows

1-10 of 37,282							Display	10	results per page
Lot Number	itk	Trial Name	Trial Site	Campaign	Rep	yield (rdt)			
Alberic	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	NA			
Alberic	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	93,4			
DI00004	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	81,8			
DI00004	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	95,1			
DI00005	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	87,6			
DI00005	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	89,7			
EM00001	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	77,8			
EM00001	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	91,8			
EM00003	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	91,1			
EM00003	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	85,4			

GnplS.Ephesis V2 Performances

- Study Subject

- No join
- Phenotype: select N+1
- Ok now
- Not for the Future

study_subject_t	
🔑 study_subject_id	BIGINT
study_subject_name	
study_subject_number	
root_stock_id	
lot_id	
trial_id	
type_id	
dn_lot_number	
dn_acc_number	
dn_acc_id	
dn_acc_name	
dn_taxon_sc_name	
dn_genus	
dn_species	
dn_subspecies	
dn_sst_type_text_code	
dn_type_name	
dn_trial_name	
dn_trial_number	
dn_site_id	
dn_site_name	
dn_levels	

phenotype_t	
🔑 phenotype_id	BIGINT
value	CHARACTER VARYING(255)
phenotyping_date	TIMESTAMP(6) WITHOUT TIME ZONE
comments	CHARACTER VARYING(255)
study_subject_id	BIGINT
observation_variable_id	BIGINT
phenotyping_campaign_id	BIGINT
pathogen_id	BIGINT
dn_obs_var_comments	CHARACTER VARYING(4000)
dn_obs_var_proto	CHARACTER VARYING(4000)
dn_obs_unit	CHARACTER VARYING(128)
dn_obs_var_ot_text_code	CHARACTER VARYING(128)
dn_obs_var_ot_name	CHARACTER VARYING(256)
dn_obs_var_ot_id	BIGINT
dn_obs_var_ot_definition	CHARACTER VARYING(4000)
dn_obs_var_ot_ontology_name	CHARACTER VARYING(256)
dn_obs_var_ot_ontology_id	BIGINT
dn_pheno_camp_name	CHARACTER VARYING(256)
dn_obs_var_ot_short_name	CHARACTER VARYING(255)

NoSQL Modelisation

- <http://blog.palo-it.com/2013/06/10/modelisation-dun-schema-dune-base-de-donnees-nosql/>
- NoSQL Document
- From Database to documents
 - Denormalisation and aggregation
 - 3 methods



Aggregations

- Document interweaving

```
{  
    "id": 10,  
    "nom": "dupont",  
    "prenom": "david",  
    "email": "me@palo-it.com",  
    "adresse":  
    {  
        "pseudo": "10 rue du test",  
        "ville": "paris",  
        "pays": "France",  
        "code postal": 75009  
    }  
}
```

- Field duplication
 - Interweaved document as attributes of root document
- Correlated documents (ie foreign Key)

```
Auteur  
{  
    "id": 10,  
    "nom": "dupont",  
    "prenom": "david",  
    "livres": [  
        101, 503, 339, 342  
    ]  
}  
  
Livre  
{  
    "id": 342,  
    "titre": "NoSQL schema",  
    "genre": "informatique",  
  
    "tags": ["informatique", "bigdata", "nosql"]  
    "auteurs": [  
        10, 234  
    ]  
}
```



Application to phenotyping

- Breeding API
 - <http://docs.brapi.apiary.io>
- Study aka Trial

```
{  
    "studyDbId": 123,  
    "studyPUI": "http://phenome-fppn.fr/phenoarch/2014/1",  
    "studyId" : "BRP-03",  
    "studyName": "Blight Resistance in Phillipines",  
    "studyObjective": "Test blight resistant cultivars",  
    "studyType": "Trial",  
    "studyLocation": "Phillipines",  
    "studyProject": "Inovine",  
    "dataSet": ["National Network", "Frost suceptibility network"],  
    "studyPlatform": "Phenome",  
    "startDate": "2015-06-01",  
    "endDate": "2015-12-31",  
    "programName": "RiceImprovementProgram",  
    "designType": "RCBD",  
    "keyContact": "Mr.PlantBreederA",  
    "contacts": [  
        {  
            "type": "scientific coordinator", "name": "John Doe", "email": "johndoe@inra.fr"  
        }]  
        "meteoStationCode": "Anlez",  
        "meteoStationNetwork": "OpenWheatherMap",  
        "studyHistory": "Previous crop was pea, then maize",  
        "studyComments": "",  
        "seasons": ["2005", "2008"],  
        "observationVariables": [  
            {  
                "observationVariableId": "CO_321:0000045",  
                "observationVariableComment": "There might be a mistake in the observation variable ID"  
            },  
            {  
                "observationVariableId": "http://www.cropontology.org/resource/CV_00000000",  
                "observationVariableComment": ""  
            }],  
        "germplasms": [  
            {  
                "germplasmDbId": "01BEL084609",  
                "germplasmPUI": "http://www.crop-diversity.org/mgis/acc/01BEL084609",  
                "germplasmName": "Pahang"  
            }]  
}
```

Application to phenotyping

- Breeding API
- Phenotypes
- One document = one data matrix line
- Elasticsearch nested documents

1-10 of 37,282 ⏪ ⏵ ⏶ | Display 10 results per page

Lot Number	itk	Trial Name	Trial Site	Campaign	Rep	yield
Alberic	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	NA
Alberic	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	93,4
D100004	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	81,8
D100004	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	95,1
D100005	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	87,6
D100005	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	89,7
EM00001	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	77,8
EM00001	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	91,8
EM00003	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	1	91,1
EM00003	t: treated	BTH_Clermont-Ferrand_2000_SetB1	Clermont-Ferrand	2000	2	85,4

```
{  
    "observationUnitDbId": 20,  
    "observationUnitPUI": "http://phenome-fppn.fr/maugio/bloc/12/23",  
    "studyId": "RIGW1",  
    "studyLocation": "Bergheim",  
    "studyPUI": "http://phenome-fppn.fr/phenoarch/2014/1",  
    "studyProject": "Inovine",  
    "studyPlatform": "Phenome",  
    "germplasmPUI": "http://inra.fr/vassal/41207Col0001E",  
    "germplasmDbId": 3425,  
    "germplasmName": "charger",  
    "treatments":  
    [  
        {  
            "factor" : "water regimen" ,  
            "modality": "water deficit"  
        }  
    ],  
    "X" : "",  
    "Y" : "",  
    "data": [  
        {  
            "observationVariableId": "CO_321:0000045",  
            "season": "2005",  
            "observationValue" : "red",  
            "observationTimeStamp": null,  
            "quality": "reliability of the observation",  
            "collectionFacilityLabel": "phenodyne",  
            "collector" : "John Doe and team"  
        },  
        {  
            "observationVariableId": "http://www.cropontology.org/cropontology.owl#OAT",  
            "season": null,  
            "observationValue" : 32,  
            "observationTimeStamp": "2006-07-03::10:00",  
            "quality": "8",  
            "collectionFacilityLabel": null,  
            "collector" : "userURIOrName"  
        }  
    ]  
}
```

GnplS,Ephesis V3 performances

- 700 000 observation units
- $3 * 10^6$ phenotypes / observations
- Elasticsearch implementation
- Response time
 - <https://urgi.versailles.inra.fr/ephesis/ephesis/viewer.do#dataResults/trials?SetIds=6,5,7>
 - 770 Trials, 49 658 documents subset, 6 variables each
 - One page, Ajax call
 - 300 – 600 ms
 - Full export
 - 23 s
- Genotyping :

- David Pilato
 - <http://david.pilato.fr/>
 - Evangelist at [elastic](#) and creator of the Elastic French Speakers [User Group](#).
- http://fr.slideshare.net/*
- Elasticsearch
 - <https://www.elastic.co/>
 - Créé en 2010
 - International
 - Applications et adoption croissante

Elasticsearch



- Lucene based
- Search engine
 - Not presented as a NoSQL database
 - NoSQL document search engine
 - BUT : it has backup systems
 - Not for long term storage
 - Aggregated documents, query oriented
 - Could be used as NoSQL DB ?
 - Storage
- HTTP, REST, JSON
- Distributed, Scalable, Cluster and Cloud ready

Cherche moi un document
de **décembre 2011** portant sur la **france**
et contenant **produit** et **david**

En SQL:

```
SELECT
    doc.*, pays.*
FROM
    doc, pays
WHERE
    doc.pays_code = pays.code AND
    doc.date_doc > to_date('2011-12', 'yyyy-mm') AND
    doc.date_doc < to_date('2012-01', 'yyyy-mm') AND
    lower(pays.libelle) = 'france' AND
    lower(doc.commentaire) LIKE '%produit%' AND
    lower(doc.commentaire) LIKE '%david%';
```

<http://fr.slideshare.net/dadoonet/elasticsearch-esme-sudria>

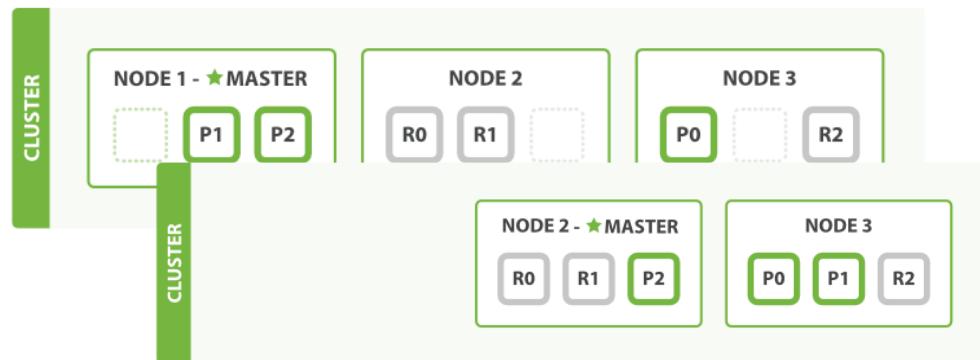
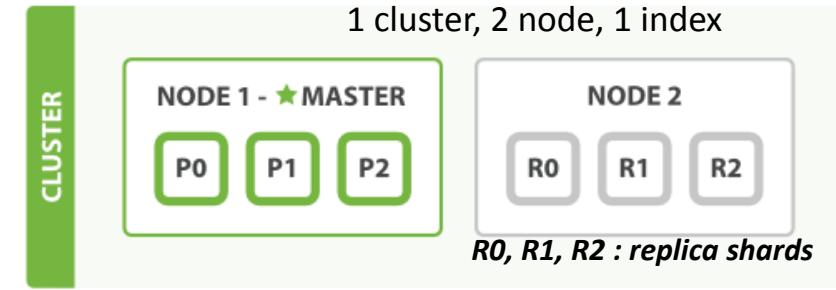
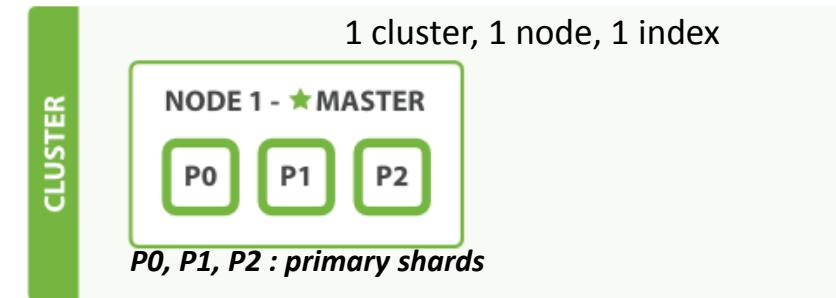


The screenshot shows a search interface with the following fields and controls:

- Power Search:** A dropdown menu labeled "Select".
- ID Number:** An input field.
- Web Title:** An input field.
- URL:** An input field.
- Category:** A dropdown menu labeled "Select".
- Web Description:** An input field.
- Keywords:** An input field.
- Contact Name:** An input field.
- Contact Email:** An input field.
- Featured Links:** A dropdown menu labeled "Select".
- Cool Links:** A dropdown menu labeled "Select".
- Bold Links:** A dropdown menu labeled "Select".
- Icon:** A section with three radio button icons: a triangle, a circle with a dot, and a square with a cross.
- Rating Average:** A dropdown menu labeled "Select".
- Number of Votes:** An input field.
- Total Hits:** An input field.
- Hits Today:** An input field.
- IP Address:** An input field.
- Submission Software Name:** An input field.

```
curl -XGET 'http://localhost:9200/docindex/doc/_search' -d '{\n    "query": {\n        "bool": {\n            "filter": {\n                "term": { "pays.libelle" : "france" },\n                "match": {\n                    "doc.commentaire": {\n                        "query": "produit david",\n                        "operator": "and"\n                    },\n                    "range": {\n                        "doc.date_doc": {\n                            "gte": "2011-12",\n                            "lt": "2012-01"\n                        }\n                    }\n                }\n            }\n        }\n    }\n}'
```

- URL HTTP :
ES_instance/index/type/documents
- node :
 - Running instance of elasticsearch, belongs to a cluster.
- cluster :
 - one or more nodes with same cluster name
 - Single master node
 - chosen automatically
 - replaced if fails.
- shard
 - single Lucene instance
 - low-level “worker”
 - managed automatically by elasticsearch.
 - primary and replica shards.

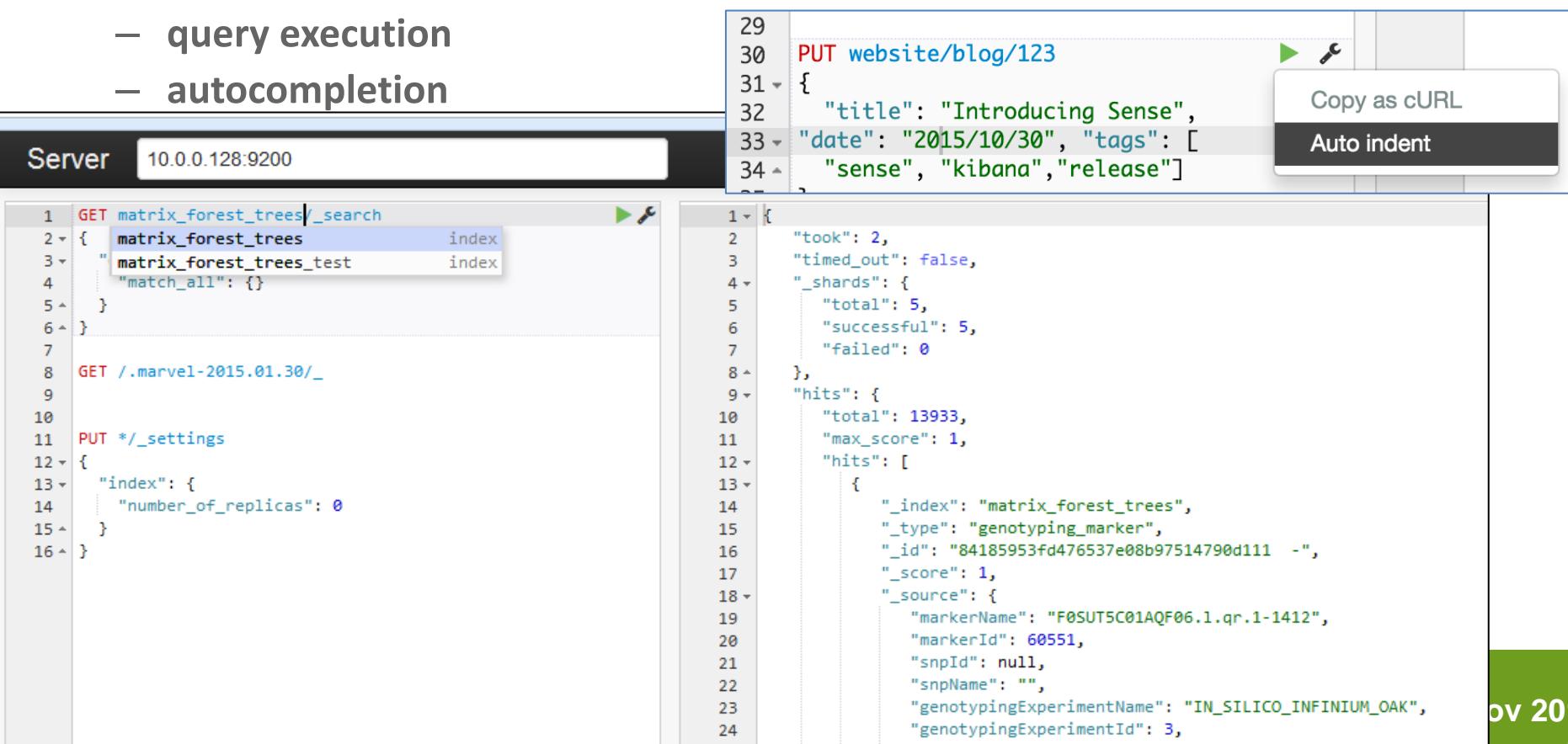


- type
 - like a *table* in a relational database.
- index
 - like a *database* in a relational database.
 - has a mappings which defines multiple types.
 - types
 - Study
 - http://localhost:9200/phenoindex/studytype/_search?
 - Phenotypes
 - http://localhost:9200/phenoindex/phenotype/_search?
- Alias
 - like a SDGB view on multiple indices (possibly filtered via a query)

Development

- Very good query DSL/API documentation
 - <https://www.elastic.co/guide/en/elasticsearch/reference/current/search.html>
- Sense
 - Free, integrated into marvel (<https://www.elastic.co/products/marvel>)
 - <https://www.elastic.co/guide/en/sense/current/sense-ui.html>
 - query execution
 - autocomplete

Server 10.0.0.128:9200



The screenshot shows the Elasticsearch Sense UI interface. At the top, there's a code editor window with the following JSON code:

```

29 PUT website/blog/123
30 {
31   "title": "Introducing Sense",
32   "date": "2015/10/30", "tags": [
33     "sense", "kibana", "release"
34   ]

```

To the right of the code editor is a results panel displaying the search results for the query "matrix_forest_trees". The results are as follows:

```

1 {
2   "took": 2,
3   "timed_out": false,
4   "_shards": {
5     "total": 5,
6     "successful": 5,
7     "failed": 0
8   },
9   "hits": {
10    "total": 13933,
11    "max_score": 1,
12    "hits": [
13      {
14        "_index": "matrix_forest_trees",
15        "_type": "genotyping_marker",
16        "_id": "84185953fd476537e08b97514790d111",
17        "_score": 1,
18        "_source": {
19          "markerName": "F0SUT5C01AQF06.1.qr.1-1412",
20          "markerId": 60551,
21          "snpId": null,
22          "snpName": "",
23          "genotypingExperimentName": "IN_SILICO_INFINIUM_OAK",
24          "genotypingExperimentId": 3,
25          "genotypingExperimentType": "in silico"
26        }
27      }
28    ]
29  }
30 }

```

On the far right of the results panel, there are two buttons: "Copy as cURL" and "Auto indent".

Aggregations, ie Facets

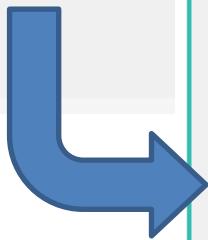
- Analytic queries
- Two main concepts:
 - **Buckets** Collections of documents that meet a criterion
 - **Metrics** Statistics calculated on the documents in a bucket

```
SELECT COUNT(color) ①  
FROM table  
GROUP BY color ②
```

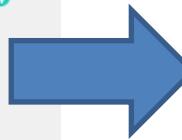
-
- ① `COUNT(color)` is equivalent to a metric.
 - ② `GROUP BY color` is equivalent to a bucket.

Aggregation example

```
GET /cars/transactions/_search?search_type=count
{
  "aggs" : { ①
    "colors" : { ②
      "terms" : {
        "field" : "color" ③
      }
    }
  }
}
```



```
{
  ...
  "hits": {
    "hits": [] ①
  },
  "aggregations": {
    "colors": { ②
      "buckets": [
        {
          "key": "red", ③
          "doc_count": 4 ④
        },
        {
          "key": "blue",
          "doc_count": 2
        },
        {
          "key": "green",
          "doc_count": 2
        }
      ]
    }
  }
}
```



- 1/ Display on result page as clickable facet
- 2/ on click on « red » filter
color : red

Aggregation : Avg and imbrication

```
GET /cars/transactions/_search?search_type=count
{
  "aggs": {
    "colors": {
      "terms": {
        "field": "color"
      },
      "aggs": {
        "avg_price": { ①
          "avg": {
            "field": "price"
          }
        },
        "make": { ②
          "terms": {
            "field": "make" ③
          }
        }
      }
    }
  }
}
```



```
{
  ...
  "aggregations": {
    "colors": {
      "buckets": [
        {
          "key": "red",
          "doc_count": 4,
          "make": { ①
            "buckets": [
              {
                "key": "honda", ②
                "doc_count": 3
              },
              {
                "key": "bmw",
                "doc_count": 1
              }
            ]
          },
          "avg_price": {
            "value": 32500 ③
          }
        },
        ...
      }
    }
  }
}
```

- Monitoring : Marvel
 - Gratuit dans ES 2.0
- Installation
 - Repository DEB and RPM
 - Unzip and run (development instance, tomcat like)
- Configuration YAML

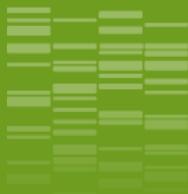
Elasticsearch architecture

- Scalability
 - Automatic index data distribution accross shards
 - Adding nodes to increase number of shards
- Shard Replication
 - Option
 - Index by index configuration
 - Data availability
 - Query performances

- Version 1.7.3
- Data :
 - 65 Gb
 - 600 millions documents, including nested
- Cluster Prod (same for dev)
 - 2 nodes: VM, 16 Go RAM, 2To, 8 CPU
- Each nodes
 - Unlimited number of indices
- Backup
 - Weekly, retention 8 weeks
 - On the fly, without service interruption, incremental
 - Via a REST query (API HTTP)
- Good data security
 - Protection against index corruption
 - Near NoSQL DB state

Insertion

- Generate JSON
 - Talend
 - Java
- Insert
 - Logstash
 - CURL



Thank you

