

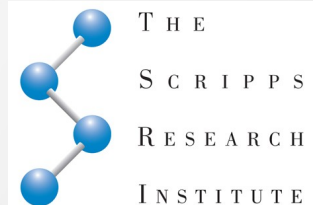
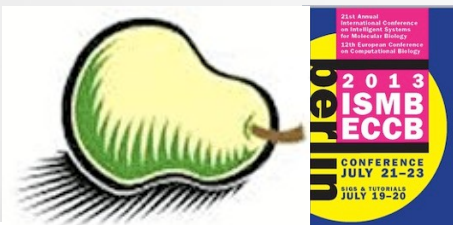
MyGene.info

- Making Elastic Gene API

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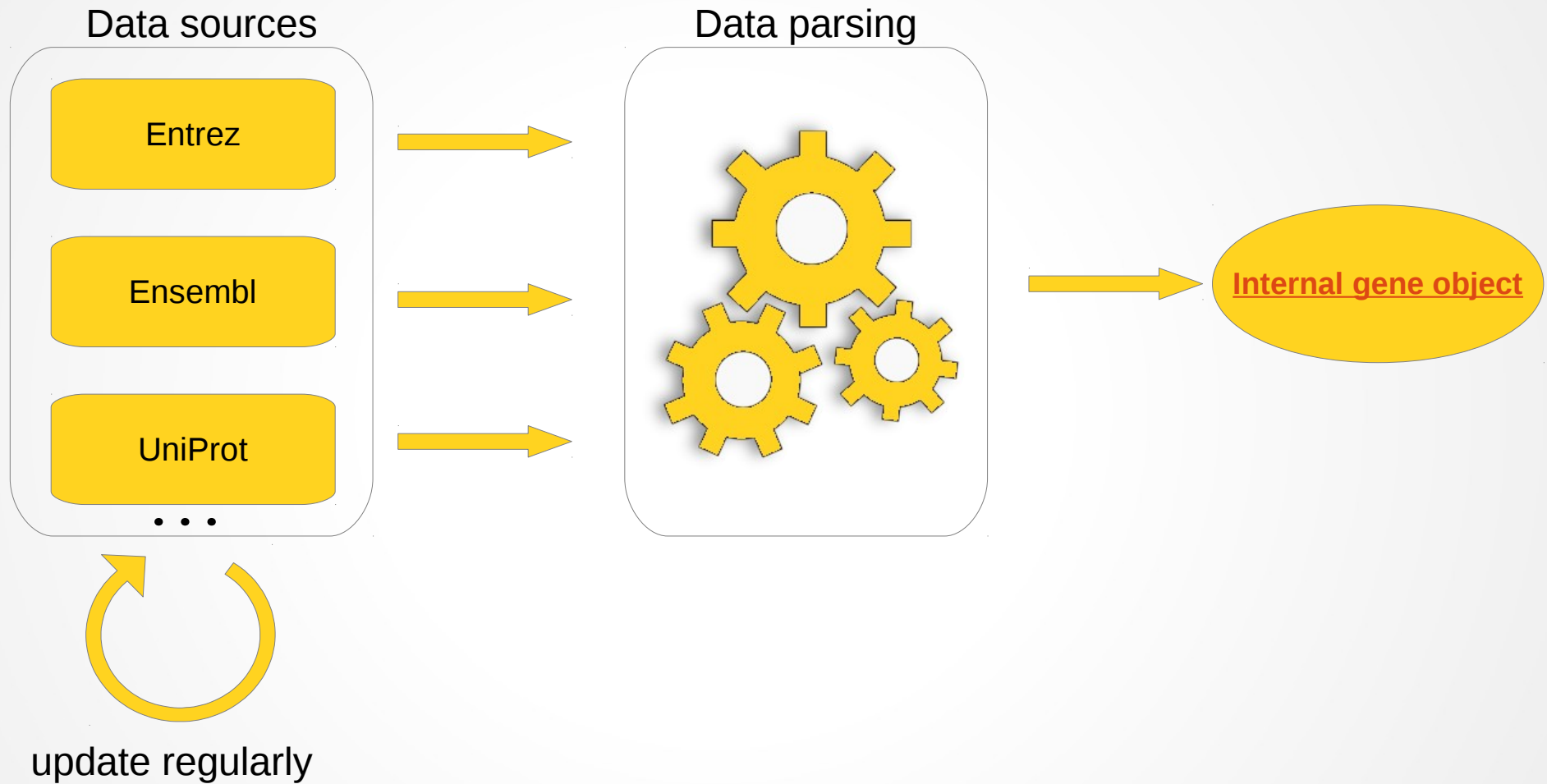
BOSC 2013
July 20, 2013



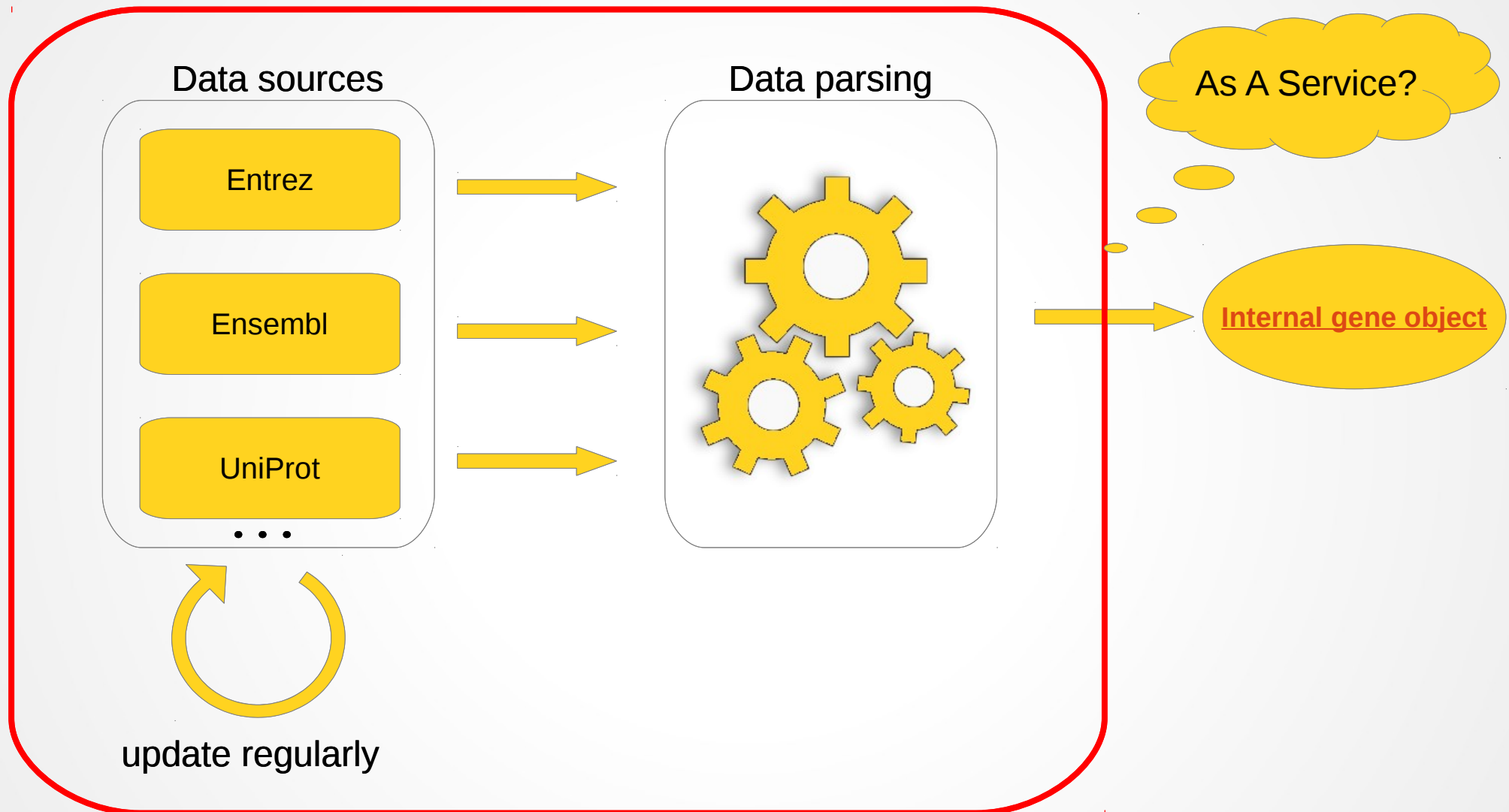
Being “Elastic”

- Fast
- Always ON
- Up-to-date
- Scalable
- Extensible

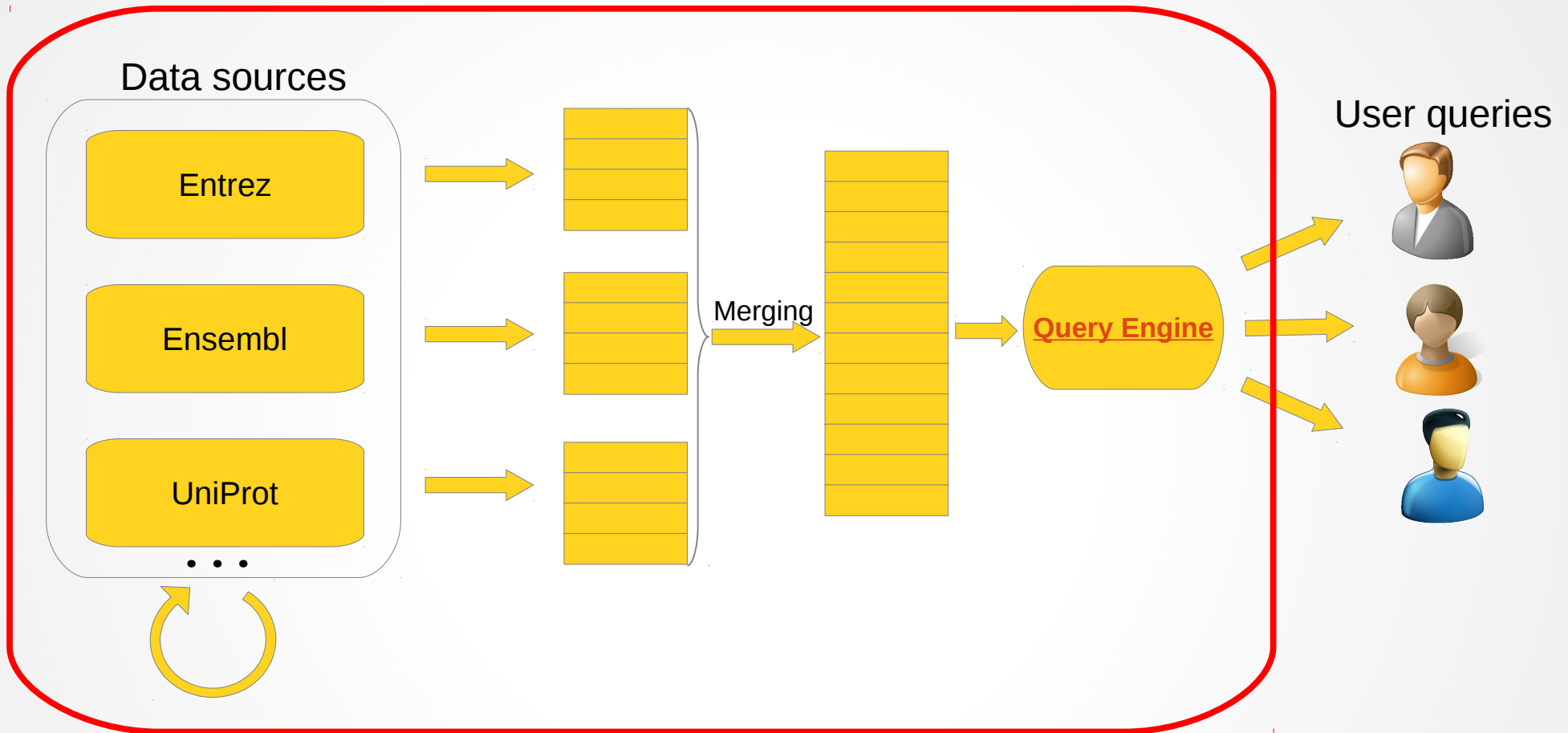
Common procedure for gene data retrieval



Common procedure for gene data retrieval

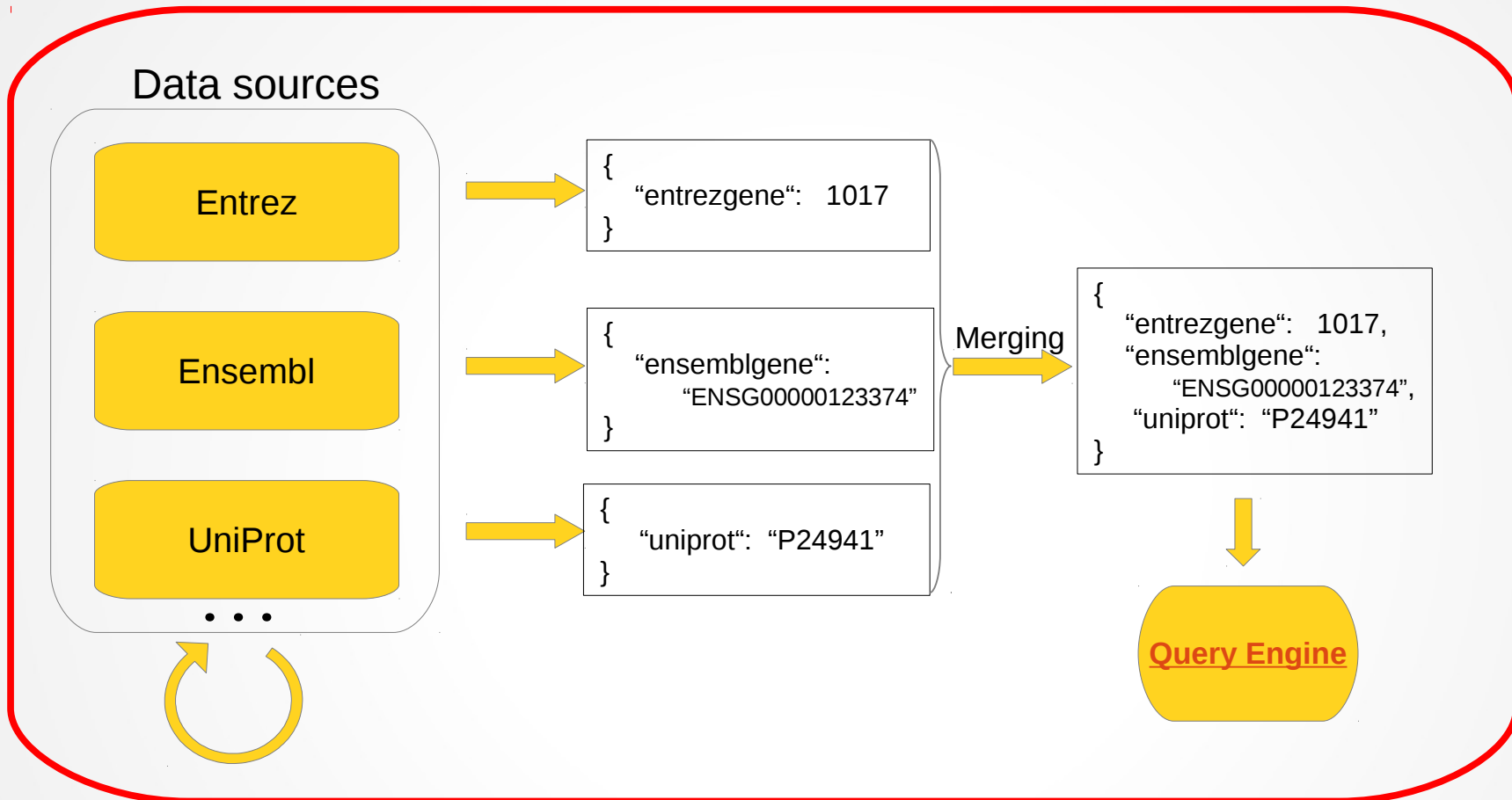


Working model - 1

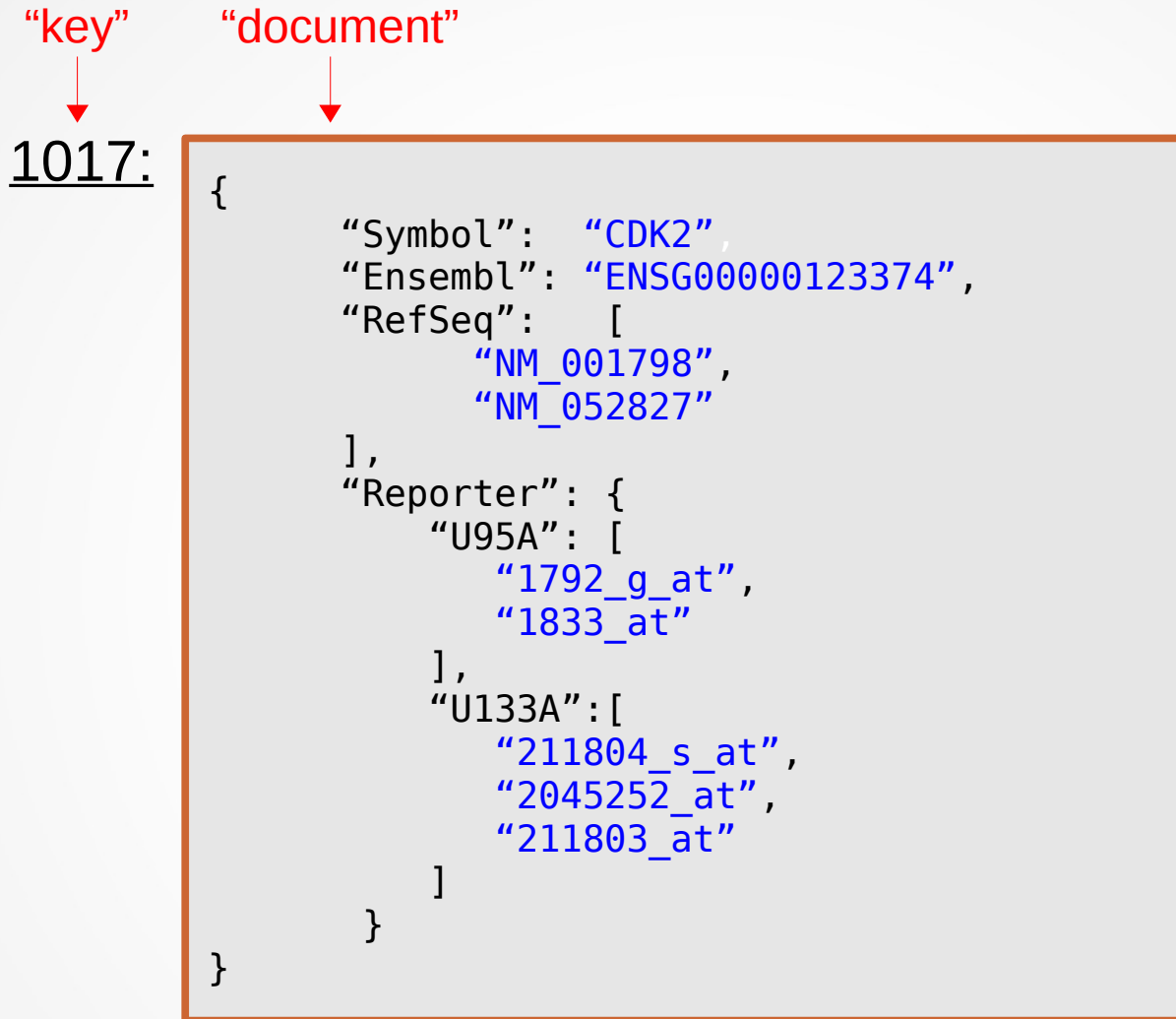


Working model - 1

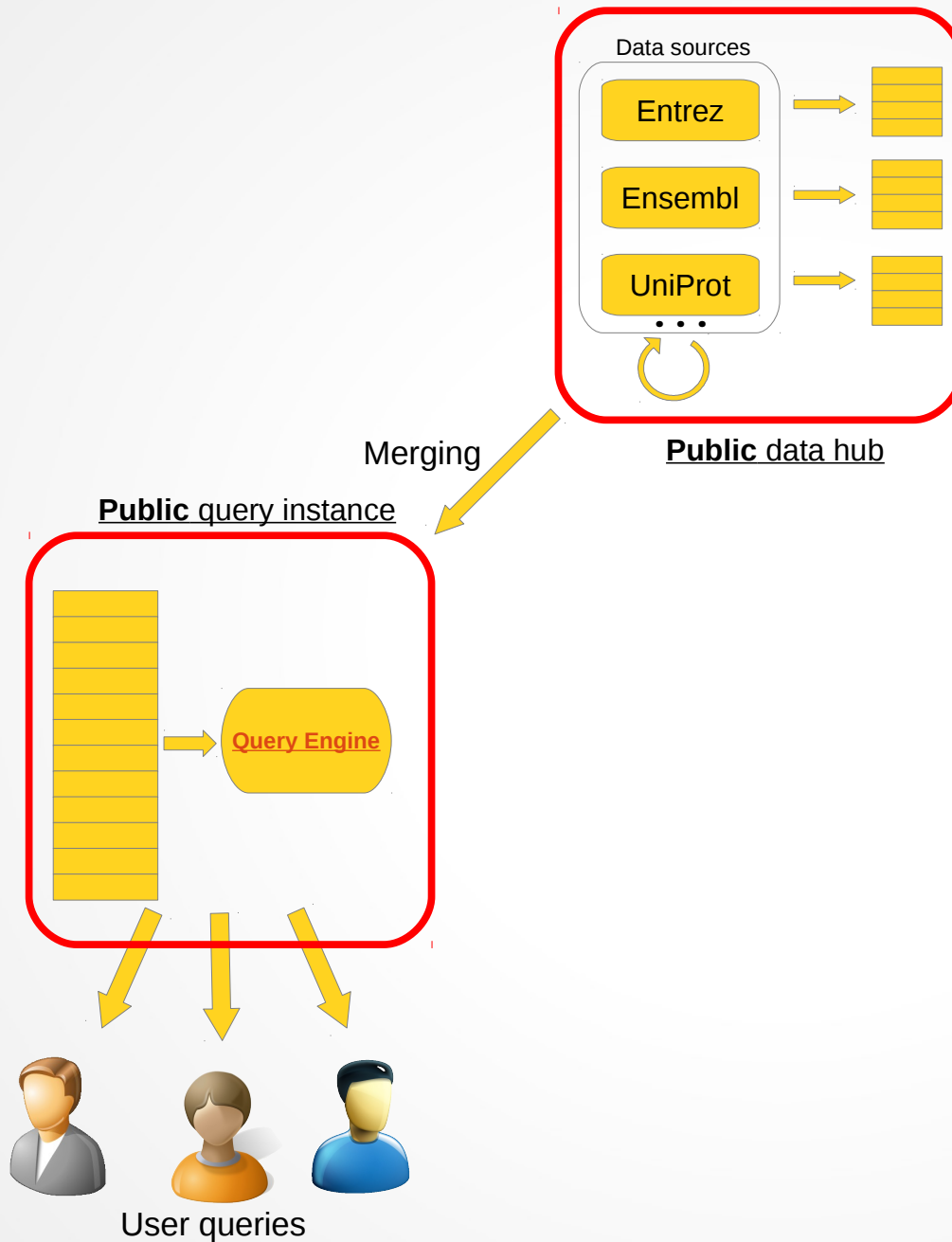
A dummy merging example:



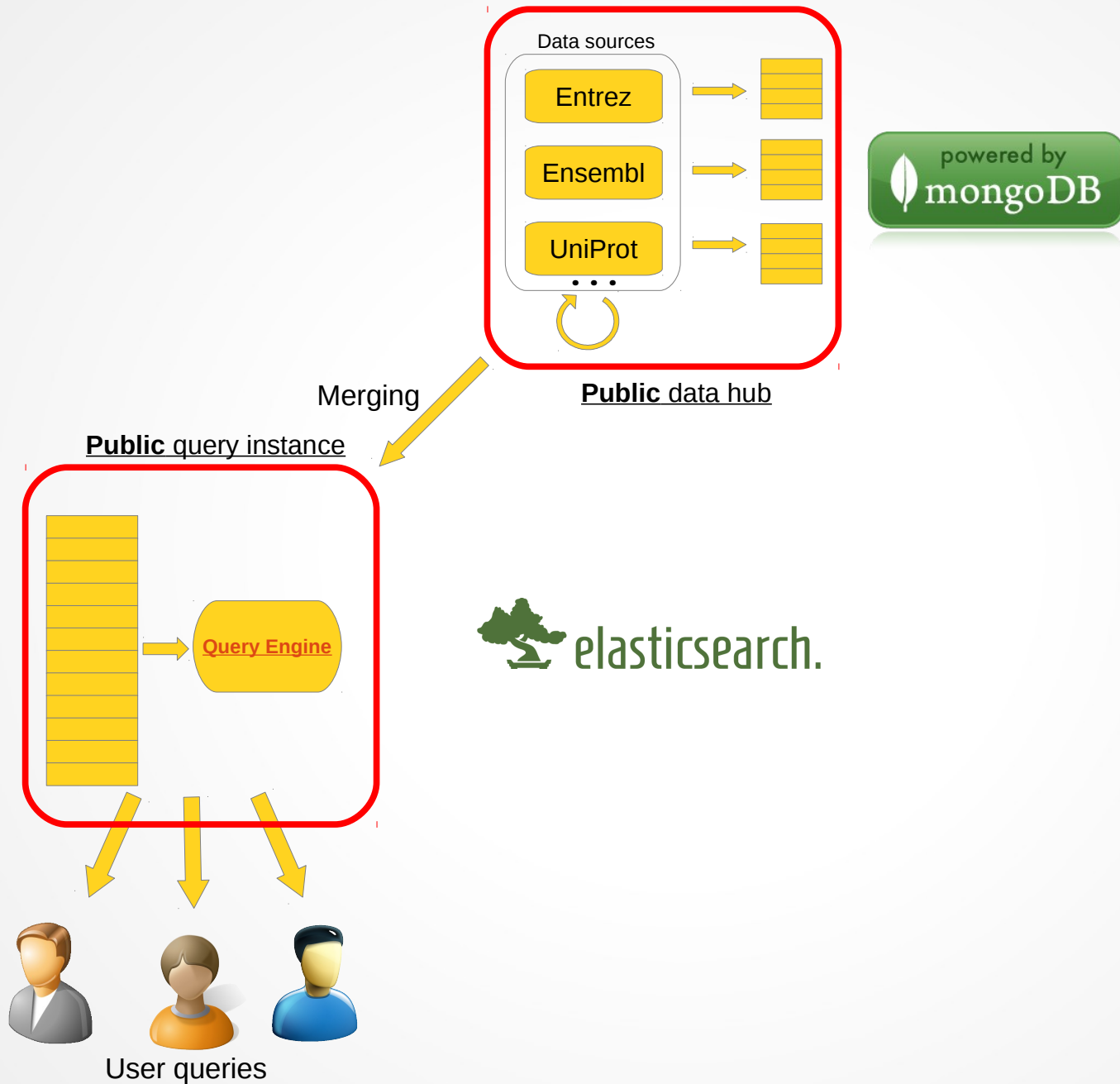
Gene object in noSQL database



Syncing from data-hub to query instance



Syncing from data-hub to query instances



Public query instance

<http://MyGene.info>

(currently v2 API, two endpoints)

any query term(s)



`http://MyGene.info/v2/query?q=<query>`



matching gene hits

gene id(s)



`http://MyGene.info/v2/gene/<geneid>`



matching gene objects

Public query instance

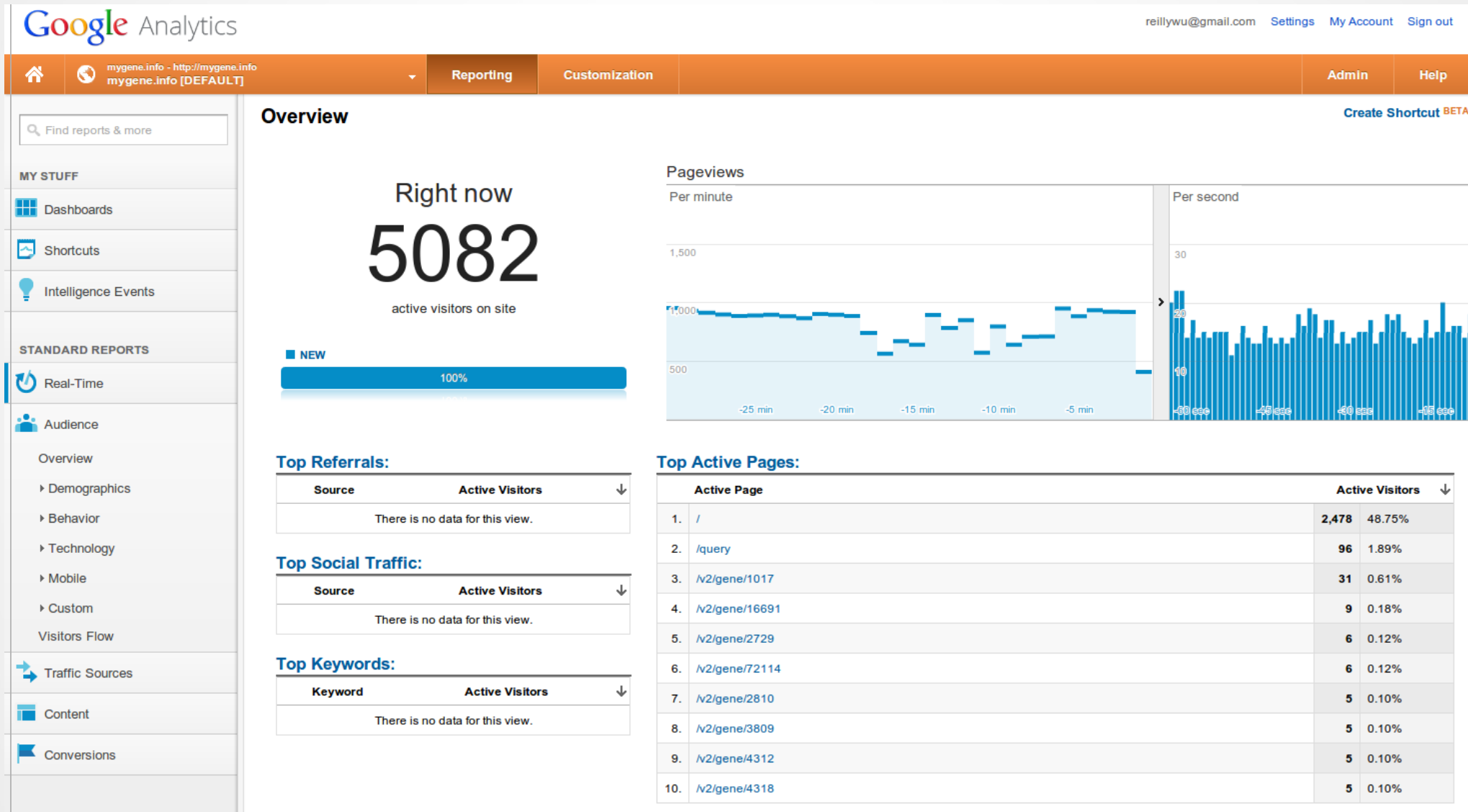
<http://MyGene.info>

- Support ALL species, from NCBI (>12K species, >13M genes)
- >40 annotation fields and expanding
- Weekly-updated
- Flexible query interface
 - Simple queries *?q=cdk2*
 - Fielded queries *?q=symbol:cdk2*
 - Wildcard queries *?q=cdk**
 - Genomic interval queries *?q=chr1:1-100,000&species=human*
 - Species filter *?q=cdk2&species=mouse,rat*
 - Returning fields filter *?q=cdk2&fields=symbol,homologene*
- Support batch queries, JSONP, CORS
- Committed for long-term availability

Public query instance

<http://MyGene.info>

High-performance host (serving ~500K requests/day)



Public query instance

Third-party packages

MyGene.py - Python wrapper

<https://pypi.python.org/pypi/mygene>

```
pip install mygene
```

```
In [1]: import mygene
In [2]: mg = mygene.MyGeneInfo()
In [3]: mg.getgene(1017)
Out[3]:
{'_id': '1017', 'entrezgene': 1017, 'name': 'cyclin-dependent kinase 2',...}
In [4]: mg.query('cdk2')
Out[4]:
{'hits': [{ '_id': '1017',
             '_score': 373.24667,
             'entrezgene': 1017,
             'name': 'cyclin-dependent kinase 2',
             'symbol': 'CDK2',
             'taxid': 9606},
          ...],
 'max_score': 373.24667,
 'took': 10,
 'total': 28}
```

Public query instance

Third-party packages

MyGene.autocomplete

- Gene query autocomplete widget

<https://bitbucket.org/sulab/mygene.autocomplete>

```
1 |$('input.mygene_query_target').genequery_autocomplete({
2   "species": "human,mouse,rat",
3   "limit": 10,
4   "gene_label": "{symbol}: {name} ({species})",
5   "select": function(event, ui) {
6     alert( ui.item ?
7       "Selected: " + ui.item.label + '('+ui.item._id+')':
8       "Nothing selected, input was " + this.value);
9   }
10 });
```

Public query instance

Third-party packages

MyGene.autocomplete

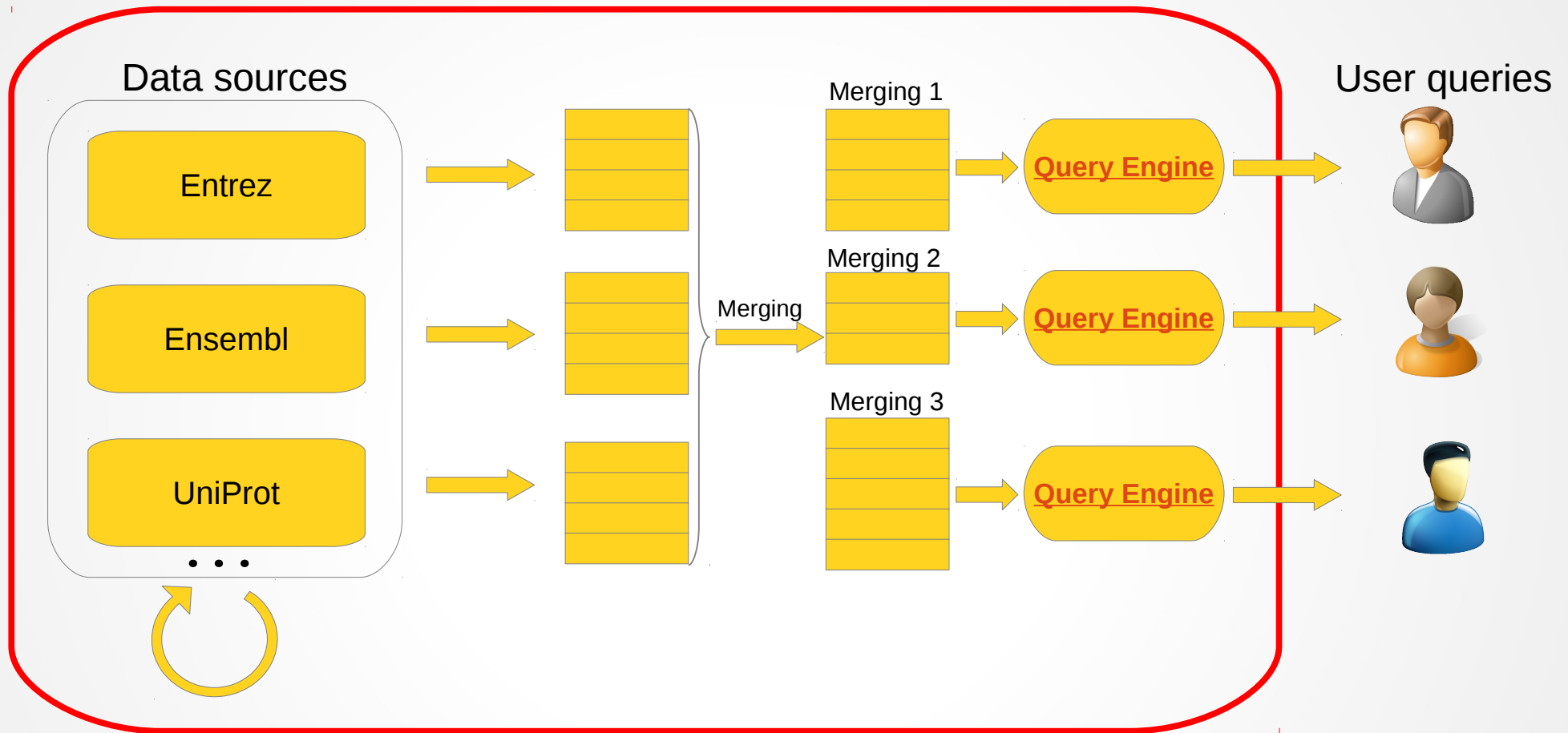
- Gene query autocomplete widget

<https://bitbucket.org/sulab/mygene.autocomplete>

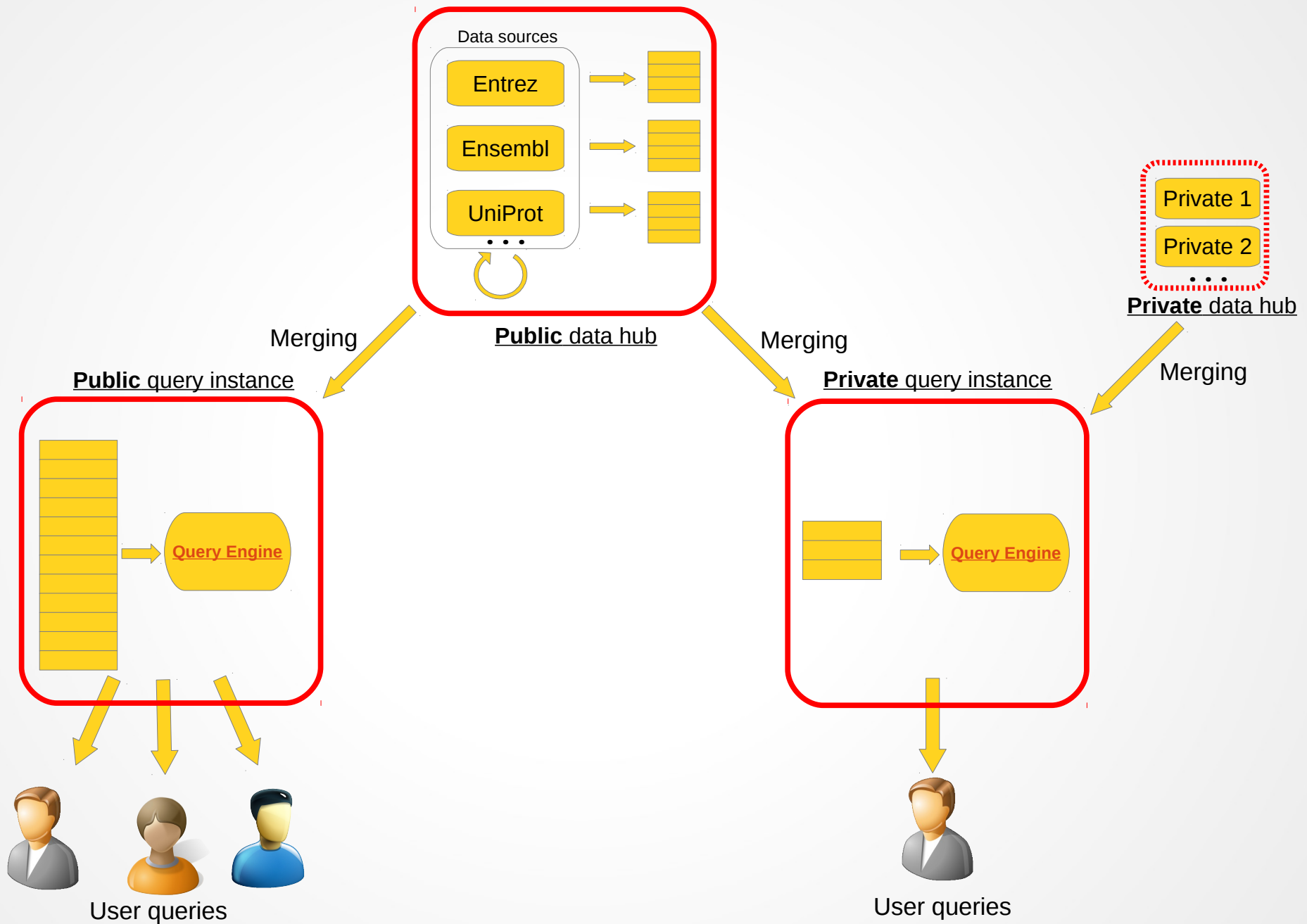
Enter a gene here:

- CDK2: cyclin-dependent kinase 2 (human)
- Cdk2: cyclin-dependent kinase 2 (mouse)
- Cdk2: cyclin dependent kinase 2 (rat)
- CDK2AP1: cyclin-dependent kinase 2 associated protein 1 (human)
- Cdk2ap1: CDK2 (cyclin-dependent kinase 2)-associated protein 1 (mouse)
- Cdk2ap2: CDK2-associated protein 2 (mouse)
- Cdk2ap1-ps5: Cdk2ap1-ps5 pseudogene (rat)
- Cdk2ap1-ps1: Cdk2ap1-pseudogene 1 (rat)
- Cdk2ap1-ps3: Cdk2ap1-ps3 pseudogene (rat)
- Cdk2ap1-ps2: Cdk2ap1-ps2 pseudogene (rat)

Working model – 2



Syncing from data-hub to query instances



Private query instance

- Dedicated host
- Same powerful query interface
 - Third-party packages still work
- Public data still get sync-ed
- Allow to merge private data

To reach us?

Questions on public query instance

or

interested in setting up your own private query instance?

Please let us know:

help@mygene.info

Code repositories

- Web front-end

<https://bitbucket.org/sulab/mygene.info>

Apache 2 licensed

- Data hub

<https://bitbucket.org/sulab/mygene.hub>

GPL v3 licensed

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