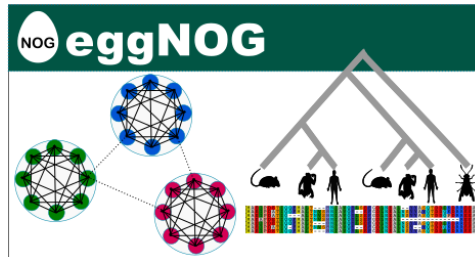


Consistently annotated
prokaryotic genomes

<http://progenomes.embl.de>

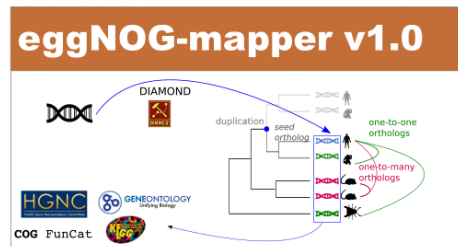
Mende et al. NAR 45 (2017) D286



Orthologous groups and
functional information

<http://egglog.embl.de>

Huerta-Cepas et al. NAR (2016)



Fast accurate functional
annotation

<http://egglog-mapper.embl.de>

Huerta-Cepas et al. MBE (2017)

ETE Toolkit

A Python framework to work with trees

```
from ete3 import Tree
tree = Tree('((A,B), D);')
```

```
print tree
#      /-A
#     /-|
#    --| \-B
#       \-D
```

```
A = tree & "A"
A.up.show()
```

Phylogenetic reconstruction
and analysis

<http://etetoolkit.org>

Huerta-Cepas et al. MBE (2016)