

# Gene Set Enrichment API – version 1.0<sup>1</sup> 28<sup>th</sup> April 2016

A simple RESTful API to analyse (Fisher-exact) enrichment of user-provided set of Ensembl gene identifiers against differentially expressed genes in each comparison in Expression Atlas.

Authors: Nuno Fonseca, Robert Petryszak

Contact: atlas-feedback@ebi.ac.uk

This document describes the RESTful API that was developed to analyse (Fisher-exact) enrichment of user provided set of Ensembl gene identifiers in a given organism against differentially expressed genes in each comparison for that organisms in Expression Atlas. The code used to perform the enrichment analysis is available from <a href="https://github.com/gxa/atlas\_gsa">https://github.com/gxa/atlas\_gsa</a> and performs Fisher-exact gene set overlap (FDR = 0.01) between the user-provided set of genes in a given organism against the set of differentially expressed genes in each Atlas comparison for that organism. All genes tested for differential expression in a given comparison are treated as background.

<sup>&</sup>lt;sup>1</sup> If you use this data in your research or service, please reference this publication. Thank You

## **Expression Atlas Organisms**

#### **Format**

Item	Description
URL	http://www.ebi.ac.uk/fg/gse/api/getOrganisms

## Gene Set Enrichment in an Organism

#### **Format**

Item	Description
URL PATTERN	ROOT/FORMAT/getOverlappingComparisons/ORGANISM/GENE_IDS
ROOT	http://www.ebi.ac.uk/fg/gse/api/
FORMAT	tsv or json
ORGANISM	An organism returned by http://www.ebi.ac.uk/fg/gse/api/getOrganisms
GENE_IDs	A list of (maximum 100) space-separated Ensembl gene identifiers in ORGANISM

## **Example Calls to Retrieve Individual Run Data**

### **URL**

ROOT/tsv/getOverlappingComparisons/arabidopsis\_thaliana/AT1G48030 AT1G53240 AT2G17130 AT2G20420 AT2G44350 AT2G47510 AT3G09810 AT3G15020 AT3G17240 AT3G27380 AT3G55410 AT3G60100 AT4G26910 AT4G35260 AT4G35650 AT4G35830 AT5G03290 AT5G08300 AT5G23250 AT5G40650 AT5G50950 AT5G55070 AT5G65165 AT5G65750 AT5G66760 ROOT/tsv/getOverlappingComparisons/homo\_sapiens/ENSG00000001167

ROOT/tsv/getOverlappingComparisons/homo\_sapiens/ENSG00000001167
ENSG00000001630 ENSG00000005339 ENSG00000012174 ENSG00000066136
ENSG00000067064 ENSG00000076555 ENSG00000079459 ENSG00000082014
ENSG00000084676 ENSG00000099194 ENSG00000100934 ENSG00000101849
ENSG00000104549 ENSG00000108424 ENSG00000109929 ENSG00000110921
ENSG00000112972 ENSG00000113161 ENSG00000113615 ENSG00000114650
ENSG00000119927 ENSG00000120837 ENSG00000125629 ENSG00000125686
ENSG00000130589 ENSG00000132142 ENSG00000132341 ENSG00000137574
ENSG00000138802 ENSG00000140396 ENSG00000140943 ENSG00000142453
ENSG00000149809 ENSG00000150961 ENSG00000152700 ENSG00000152904
ENSG00000160285 ENSG00000160752 ENSG00000163344 ENSG00000167508
ENSG00000169710 ENSG00000170522 ENSG00000172893 ENSG00000176986
ENSG00000177200 ENSG00000177565 ENSG00000188786 ENSG00000186350
ENSG00000186480 ENSG00000186951 ENSG00000188786 ENSG00000198646



## **Returned Fields**

Field	Description
EXPERIMENT	Expression Atlas experiment accession
COMPARISON_ID	Comparison identifier in EXPERIMENT
P-VALUE	P-value returned by the Fisher-exact test
OBSERVED	The number of user-provided genes that are differentially expressed in a given comparison
EXPECTED	The number of user-provided genes that have been tested in a given comparison
ADJUSTED P-VALUE	P-value adjusted by the multiple testing correction
EFFECT SIZE	OBSERVED/EXPECTED
COMPARISON_TITLE	The title of the COMPARISON_ID in EXPERIMENT
EXPERIMENT_URL	URL for COMPARISON_ID in EXPERIMENT

# **Acknowledgements**

The work described above is funded by the BBSRC, for which we express our gratitude. We would also like to thank all members of the Gene Expression Team without whom none of this would have been possible.

