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Find Proteins of Unknown Function (PUFs) using Plantannot -Protocol B

Forked from Find Proteins of Unknown Function (PUFs) using Plantannot - Protocol A

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1FMRRAPA

1	Works for me	dx.doi.org/10.17504/protocols.io.bgdgjs3w
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ABSTRACT

The Plantannot software provides several filters and a text search box that allows searching for molecules by its desired annotation features. These filters are needed to obtain PUFs and to try to relate them to abiotic stresses using RNA-seq expression data and co-expression networks. The Filters menu is separated in 8 fields, of those we are going to use only five: "Organism", "Feature type", "Orthology", "Orthologs_coexpression" and "Analyses". The "Feature Type" filter has three molecule types, from those the polypeptide box is the only that is going to be always checked and the others blank. By using the other 4 remaining filters, 6 protocols were created as examples of different ways to selecting PUFs. Protocol A: using lack of both homology and protein domain signatures. Protocol B: using lack of homology, presence of domain signatures - trying to select Domains of Unknown Function (DUF) from PFAM, and the text search "Unknown function". Protocol D-F: same protocols of A-C but using ortholog groups to find homolog proteins with co-expression data related to abiotic stress.

Protocol B is intended to find PUFs from organisms whose proteins are not yet in the NCBI's "nr" database and have DUF domains from PFAM found by InterproScan, selected using the text search "Unknown function".

EXTERNAL LINK

https://www.machado.cnptia.embrapa.br/plantannot

Entering application

1 Enter the Plantannot Result's page, with empty filters and text box search: https://www.machado.cnptia.embrapa.br/plantannot/find/?q=

Or you can enter the https://www.machado.cnptia.embrapa.br/plantannot initial page and click on the magnifying glass with the text box empty as well.



https://www.machado.cnptia.embrapa.br/plantannot

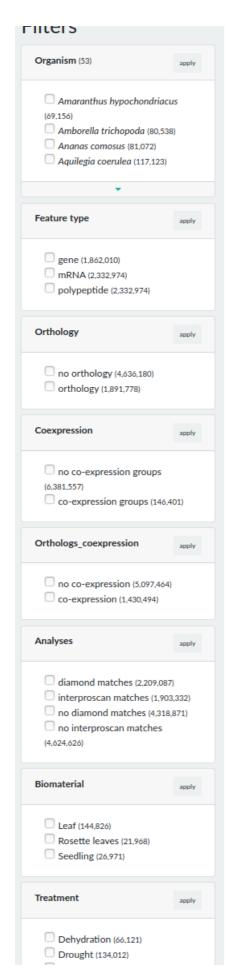
Filtering

2 Find PUFs from organisms whose proteins are not yet in the NCBI's "nr" database and have PFAM's DUF domains found by InterproScan. Proteins will be selected using the text search "Unknown function".

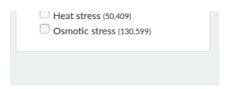
Visualize the "Filters" card on the left of the page from step 1:

Eiltore

Citation: Marcos Viana, Mauricio Mudadu, Adhemar Zerlotini (05/20/2020). Find Proteins of Unknown Function (PUFs) using Plantannot - Protocol B. https://dx.doi.org/10.17504/protocols.io.bgdgis3w



 $\textbf{Citation:} \ \ \text{Marcos Viana, Mauricio Mudadu, Adhemar Zerlotini (05/20/2020)}. \ \ \text{Find Proteins of Unknown Function (PUFs) using Plantannot - Protocol B.} \\ \underline{\text{https://dx.doi.org/10.17504/protocols.io.bgdqjs3w}}$



https://www.machado.cnptia.embrapa.br/plantannot/find/?q=

2.1 In the "Organisms" filter, select any organisms (expand the organism's list using the green arrow) or select all by leaving all boxes empty. We will use *Oropetium tomaeum* as example. Click "apply" to execute the filter:

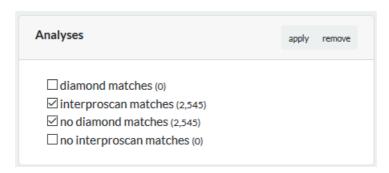
Organism (53)	apply
Amaranthus hypochondriacus	5
(69,156)	
Amborella trichopoda (80,538)	
Ananas comosus (81,072)	
Aquilegia coerulea (117,123)	
Arabidopsis halleri (78,830) Arabidopsis lyrata (97,337)	
Arabidopsis thaliana (98,188)	
Boea hygrometrica (143,334)	
Boechera stricta (87,040)	
Brachypodium distachyon	
(140,254)	
Brachypodium stacei (102,612))
Brassica oleracea (106,200)	
Brassica rapa (127,232)	
Capsella grandiflora (77,927)	
Capsella rubella (83,415)	
Carica papaya (83,355)	
Citrus clementina (92,391)	
Citrus sinensis (117,673)	
Cucumis sativus (82,231) Daucus carota (96,349)	
Eucalyptus grandis (128,909)	
Eutrema salsugineum (84,919)	
Fragaria vesca (98,493)	
Glycine max (233,338)	
Gossypium raimondii (192,039))
Kalanchoe fedtschenkoi (121,3	
Kalanchoe laxiflora (188,815)	
Linum usitatissimum (130,439)	
Malus domestica (190,548)	
Manihot esculenta (115,795)	
Medicago truncatula (175,532) Mimulus guttatus (95,286))
Musa acuminata (109,584)	
Oropetium thomaeum (85,338)
Oryza sativa (147,037)	,
Panicum hallii (136,936)	
Panicum virgatum (348,885)	
Phaseolus vulgaris (101,423)	
Populus trichocarpa (187,361)	
Prunus persica (121,051)	
Ricinus communis (93,663)	
Salix purpurea (160,905)	
Setaria italica (120,586)	
Setaria viridis (132,402)	
Solanum lycopersicum (104,17	(5)
Solanum tuberosum (151,458) Sorghum bicolor (128,371)	
Spirodela polyrhiza (58,869)	
Theobroma cacao (118,260)	
Trifolium pratense (122,552)	
Vitis vinifera (79,038)	
Zea mays (241,000)	
Zostera marina (61,350)	

 2.2 In the "Feature type" filter, select "polypeptide", and click "apply" to execute the filter:

Feature type	apply	remove
☐ gene (o) ☐ mRNA (o) ☑ polypeptide (2,551)		

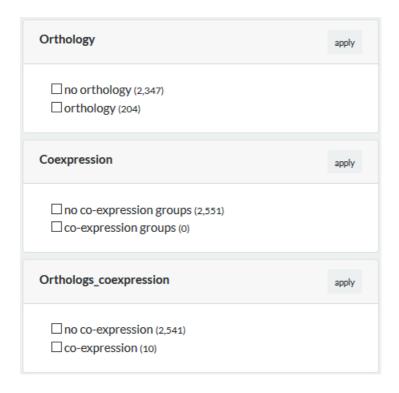
 $\frac{https://www.machado.cnptia.embrapa.br/plantannot/find/?}{q=\&selected_facets=organism%3AOropetium+thomaeum\&selected_facets=so_term%3Apolypeptide}$

2.3 In the "Analyses" filter, select both "no diamond matches" and "interproscan matches", and click "apply" to execute the filter:

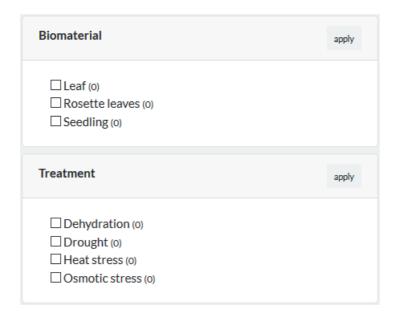


 $\frac{\text{https://www.machado.cnptia.embrapa.br/plantannot/find/?}}{\text{g=\&selected_facets=organism%3AOropetium+thomaeum\&selected_facets=so_term%3Apolypeptide\&selected_facets=analyses%3Ainterproscan+matches&selected_facets=analyses%3Ano+diamond+matcheselected_facets=analyses%3Ano+diamond+mat$

2.4 Leave the "Orthology" and "Coexpression" and "Orthologs_coexpression" filters empty:



2.5 Leave the "Biomaterial" and "Treatment" filters empty:



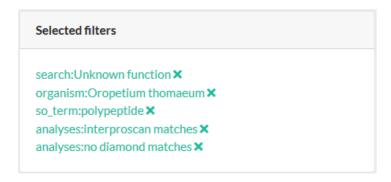
2.6 In addition to the applied filters, text search for "Unknown function" in the search box. After entering the text click on the magnifying glass icon to apply the search.



 $\frac{\text{https://www.machado.cnptia.embrapa.br/plantannot/find/?}}{\text{g=Unknown+function\&selected_facets=organism%3AOropetium+thomaeum\&selected_facets=so_ter}}$ $\frac{\text{m%3Apolypeptide\&selected_facets=analyses%3Ainterproscan+matches\&selected_facets=analyses%3Ainterproscan+matches}}{\text{Ano+diamond+matches}}}$

Filters

3 After execution of all filters we will have the following list of filters:



 $\frac{\text{https://www.machado.cnptia.embrapa.br/plantannot/find/?}}{\text{q=Unknown+function\&selected_facets=organism%3AOropetium+thomaeum\&selected_facets=so_term%3Apolypeptideselected_facets=analyses%3Ainterproscan+matches&selected_facets=analyses%3Ano+diamond+matches}}$

Viewing results

4 Visualize the "Results" card on the center-right of the screen, we will have the resulting list of *Oropetium's* PUFs, 8 PUFs were filtered:

