

# Comparison of cluster finder algorithms on cosmoDC2 data

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# Outline

#### **Cosmology from cluster count**



- Scope of the project
- Cluster catalog status
- Performance comparison
- redMaPPer/WaZP comparison

# Scope of the project

DESC project 248 (link)



- Comparison on both cosmoDC2 and DC2 datasets
- Use of CLEVAR for catalog matching and for performance estimation (common framework, try to automatize)
- Algorithms: redMaPPer, WaZP and AMICO

#### Part 1. Overall comparison

- Completeness and purity
- Mass-richness scaling relation
- Cluster redshift coverage and accuracy

#### Part 2. Direct (1-to-1) comparisons between catalogs

- Matched/unmatched clusters
- Properties for matched clusters:

richness, redshift, centering, etc.

Several DESC friend projects

# **Euclid Cluster Finder Challenge**

Euclid preparation III. Galaxy cluster detection in the wide photometric survey, performance and algorithm selection [arXiv:1906.04707]



Goal: select the best-for-Euclid-mission cluster detection algorithms 6 detection algorithms considered  $\rightarrow$  2 selected (AMICO and PZWav)

# Catalog status

Sky areas: cosmoDC2: 440 deg<sup>2</sup> cosmoDC2 small: 57 deg<sup>2</sup> DC2: 303 deg<sup>2</sup>

redMaPPer

- cosmoDC2

- DR6 catalog

 $\Lambda > 5$ 



Run 6685



# CLEVAR

### DESC package developed by M. Aguena (as DESC pipeline scientist)

# **Cluster Evaluation Resources (ClEvaR)**

© Build and Check passing coverage 100% Library to validate cluster detection. A detailed documentation of the code can be found at https://lsstdesc.org/clevar.

#### https://github.com/LSSTDESC/clevar

Matching between catalogs (proximity or membership)

**Recovery rates** 

Scaling relations (e.g mass-richness)



**API Documentation** 

#### Recovery rate

Compute recovery rates, they are computed in mass and redshift bins. There are several ways they can be displayed: - Single panel with multiple lines - Multiple panels - 2D color map

from clevar.match\_metrics import recovery

#### Simple plot

The recovery rates are shown as a function of redshift in mass bins. They can be displayed as a continuous line or with steps:

zbins = np.linspace(0, 2, 21)
mbins = np.logspace(13, 14, 5)
info = recovery.plot(c1, 'cross', zbins, mbins, shape='line')
plt.show()



They can also be transposed to be shown as a function of mass in redshift bins.

# Putting tools in place

#### Dedicated package, heavily using CLEVAR https://github.com/LSSTDESC/cluster\_challenge



Complex task, several procedures possible



Adapted from arXiv:0903.3411

Matching performed with CLEVAR proximity: 'delta\_z':0.05, 'match\_radius': '1 mpc' cross-matched pairs selected

### Completeness

#### Completeness (m,z) = N\_cl\_cross\_matched / N\_halos (m,z)



## Purity

#### Purity (r,z) = N\_halo\_cross\_matched / N\_cl (r,z)







Richness definitions are different

High purity even at very low richness: to be checked...

# redMaPPer/WaZP direct comparison (1/2)



#### After NGALS>25 and $\lambda$ >20 cuts

	N_all	N_cross	N_excl
WaZP	668	380 (57%)	288 (43%)
redMaPPer	536	380 (71%)	156 (29%)

NGALS>25  $\leftrightarrow \lambda$ >20 (similar to DES Y1 analysis)

# redMaPPer/WaZP direct comparison (2/2)

#### Stage de Franck Lesplingart (M1) en cours au LAPP



#### Color-magnitude diagrams (cross-matched pairs, $z_{cl}$ >0.8)



# Conclusion

- Cluster comparison project now active
- → Catalogs being prepared and tools being put in place
- Mass-richness studies just started
- Determine selection functions (completeness, purity and massrichness relation)
- Analyse DC2 catalogs
- Goal: publication around Summer 2023