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The Voronoi-Delauney algorithm for groups reconstruction (Marinoni et al. 2002, Gerke et al. 2005)

The algorithm computes the **Voronoi-Delaunay mesh** and then the search is divided in 3 steps:

Phase 1 - 3-D identification of groups seeds
 Phase 2 - Determining clustering strenght
 Phase 3 - Adaptive scaling based on phase-2
 richness

Phase 1: identification of groups seeds

A cylinder of radius **R1** and half-length **L1** is centered on each galaxy:

-R1 and L1 are input parameters

 - all 1st order Delauney-connected galaxies inside the cylinder are called **"1 first order neighbours"** : these galaxies + central galaxy form a **seed**;

- if there are no Delauney-connected galaxies inside the cylinder, the galaxy is considered **isolated**.

 \rightarrow R1 and L1 determine the number of seeds

Phase 2: Determining clustering strenght

A cylinder of radius **R2** and half-length **L2** is centered on the **phase-1 centre of mass**:

- R2 and L2 are input parameters

all galaxies that are Delauney-connected to first order neighbours AND that are inside this cylinder are called **"second order neighbours"** : central galaxy + 1st order
+ 2nd order neighbours are the **core** of the group;

→(central galaxy + #1st + #2nd) = group central richness N_{II}

→ N_{II} has to be corrected as a function of redshift because of the flux limit of the sample:

$$N_{II}^{corr} = \left[\frac{\langle v(z) \rangle}{\langle v(zref) \rangle}\right]^{-1} N_{II}$$

→ <v(z)> is the mean comoving number density at any given redshift z
→ zref is a zero-point redshift

Phase 3: Determining cluster dimensions and members

A cylinder of radius **R3** and half-length **L3** is centered on the **phase-2 centre of mass**:

- **R3** and **L3** depend on N_{II}^{corr} :

$$R3 = r(N_{II}^{corr})^{1/3} \qquad L3 = l(N_{II}^{corr})^{1/3}$$

where r and I are input parameters

 - all galaxies not yet processed that are inside this cylinder are considered to be members of the group;

→(central galaxy + #1st + #2nd + #phase-3 new) = group members

Testing the VDM algorithm to VVDS-like mocks (33% sampling rate, 275 km/s redshift error, $I_{AB} \leq 24$)

- **Completeness C** of a group catalogue: fraction of real groups that are successfully identified in the reconstructed catalogue

- **Purity P** of a group catalogue: fraction of reconstructed groups that correspond to real groups

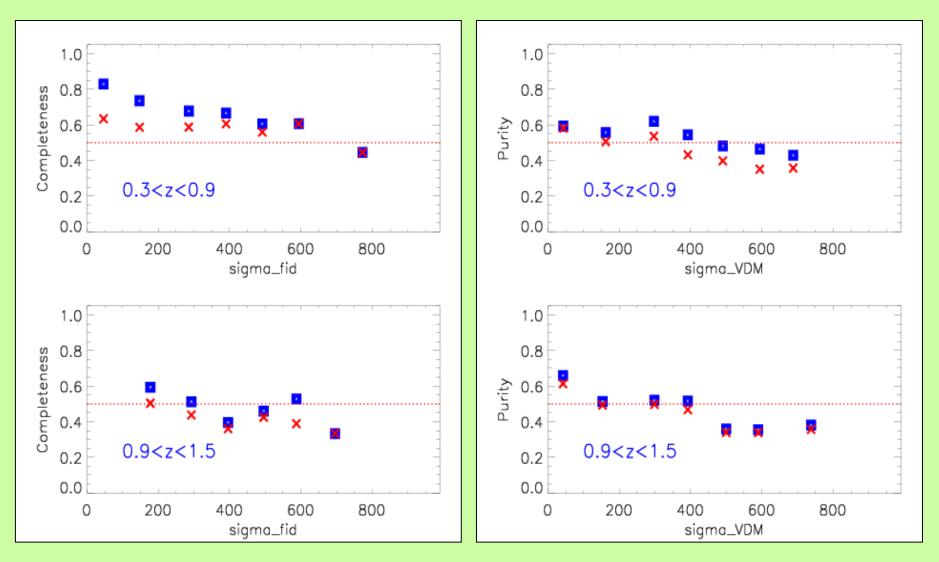
→ Trial and error to identify the best set of parameters that maximize C, keeping P always above 0.5

Testing the VDM algorithm to VVDS-like mocks (33% sampling rate, 275 km/s redshift error, $I_{AB} \leq 24$)

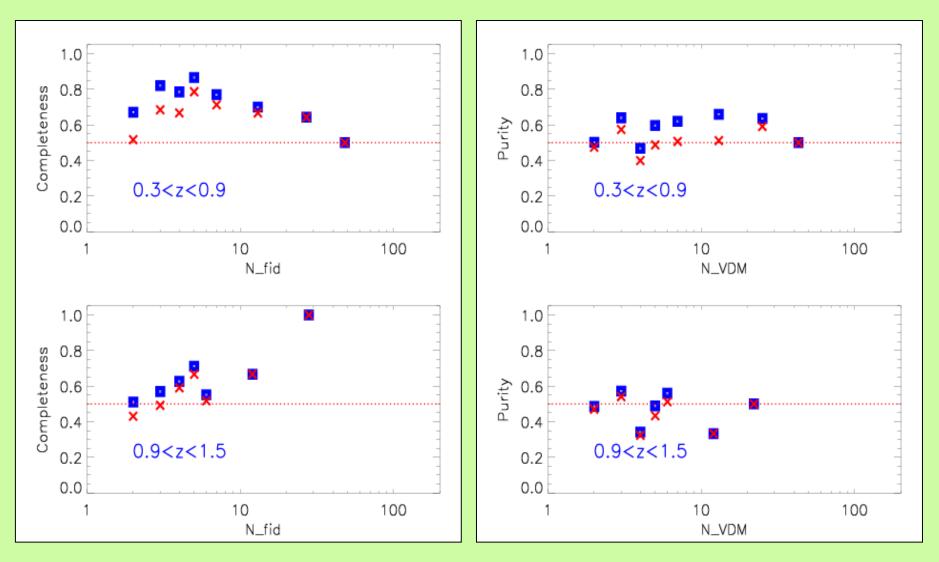
Best set (h ⁻¹ Mpc)					
R1	0.3	R2	0.5	r	0.55
L1	5.0	L2	6.0	1	15

C1 = 79.2 % P1 = 55.6 % C2 = 72.0%P2 = 49.4%

Testing the VDM algorithm to VVDS-like mocks $[R1=0.3 \quad L1=5.0 \quad R2=0.5 \quad L2=6.0 \quad r=0.55 \quad I=15.0]$



Testing the VDM algorithm to VVDS-like mocks $[R1=0.3 \quad L1=5.0 \quad R2=0.5 \quad L2=6.0 \quad r=0.55 \quad I=15.0]$



Density contrast field

Mean intergalactic separation

