



Directional-iDBSCAN

first look at LIME data

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Last presentation

- A quantitative analysis inspired by the iDBSCAN article (⁵⁵Fe based analysis) was done using the iDDBSCAN algorithm.
 - DOI: 10.1088/1748-0221/15/12/T12003
 - Selected runs: 2054 (EN), 2156 (NRAD) and 2163 (ER)
 - Link of the presentation: here

• The iDDBSCAN was able to reduce background noise in the low energy region (by reconstructing short, medium and long tracks).

• It was agreed in that meeting that the next step would be to analyse the algorithm with LIME data.



LIME data

• The first look at LIME data was done with the same parameters used in LEMOn data.

- The chosen runs:
 - [3737-3791]: Ambe runs (focus on 3790).
 - [3792-3794]: Cosmics runs soon after source off (focus on 3793).
 - This set of data was affected by the light entering the camera corners (sigma used was 1.6).

Run 3790



Clusters found DBSCAN



Event 711

Event 712









(should be avoided)

Solutions

- Use the expand_noncore parameter as *False* might solve the noise problem.
 - It is advisable to use it as *True* for a better directional clustering, especially when the tracks are faint.

- Tune the parameters to the LIME data.
 - dir_minsamples may be raised so small clusters won't be tested in the DBSCAN seeding.

• Improve the DBSCAN seeding selection.

DBSCAN seeding

- The clusters found by the iDBSCAN are seeds.
 - Each seed is tested with the RANSAC if it has more points than the dir_minsamples parameter.
 - The seeds with a good RANSAC model (the accuracy is compared with the dir_min_accuracy parameter) have one of their core points saved for the directional clustering.

• The solution is to change the RANSAC to the one used before Emanuele's implementation in *"lime21"*. (7 months ago)

RANSAC - DBSCAN seeding

- RANSAC (scikit-learn library):
 - min_samples = 50% of the set.
 - o residual_threshold = median(y median(y)).
 - The residual_threshold should be tested (if it is equal to 0 the function raises an error).
- It has difficulties to select horizontal/vertical cosmic clusters and is good to reject circular clusters.
 - The residual_threshold calculated by the function is pretty small on horizontal and circular tracks.
- Changes needed to select horizontal/vertical tracks while reject circular tracks:
 - Set a minimum value of residual threshold.
 - Rotate the clusters that did not have good RANSAC fits and test again.



Conclusions

• The two modifications worked well to reject circular tracks on the DBSCAN seeding.

• The second modification (rotate clusters) is better at selecting horizontal/vertical tracks.

- Performance at highly curved tracks still to be tested.
 - Possible modification is to use RANSAC with polynomial regression and keep using the standard residual_threshold.

Next steps

• Do speed tests to choose the fastest modification to be implemented.

- Tune the iDDBSCAN parameters in order to do a quantitative analysis.
 - Which dataset should be used?

• Any ideas or suggestions?