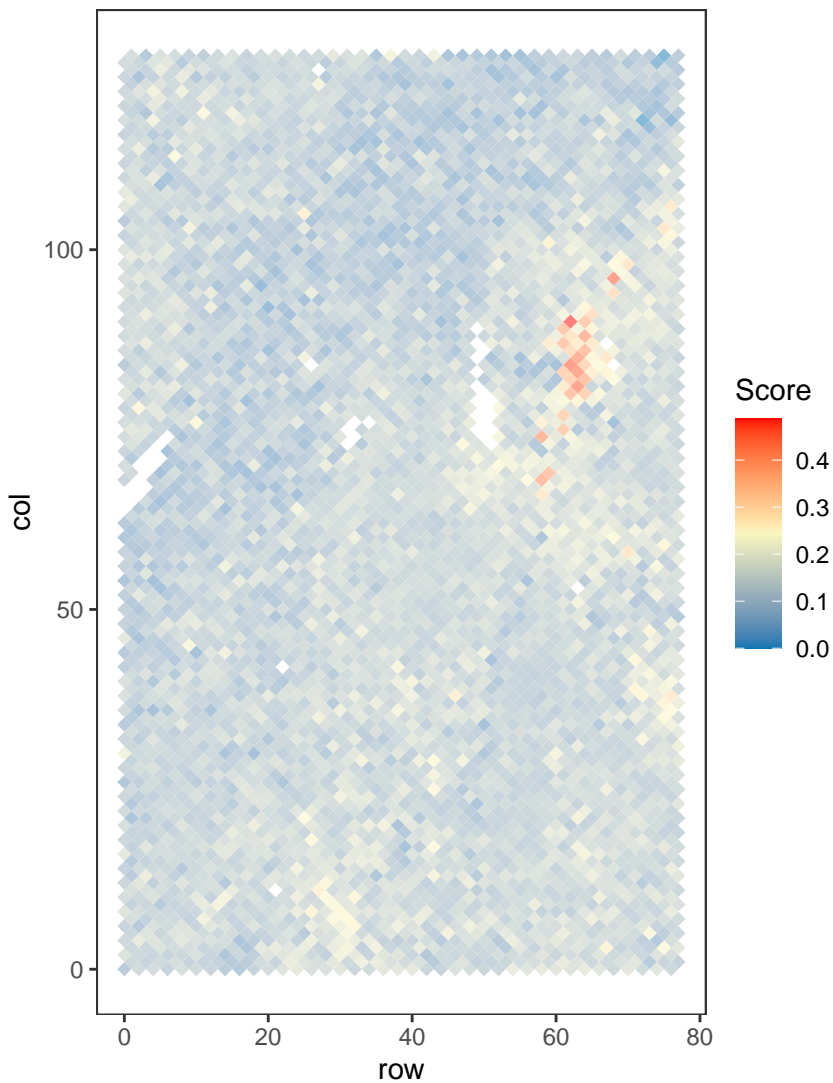


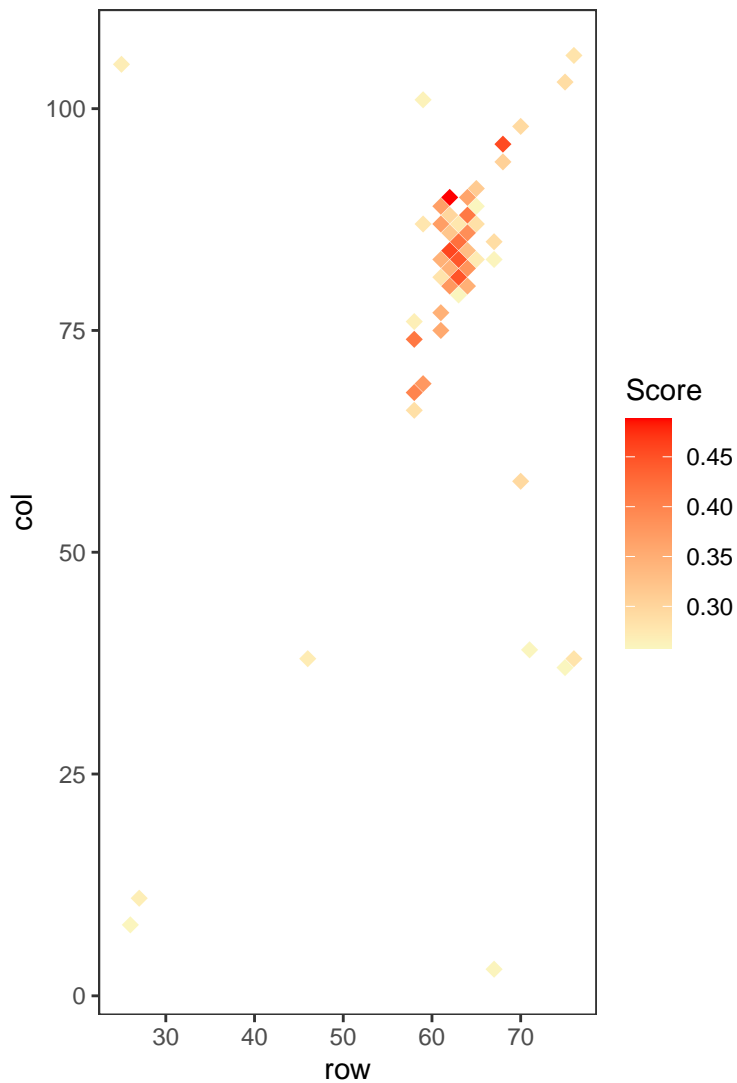
highlight

- Group_1
- Unselected

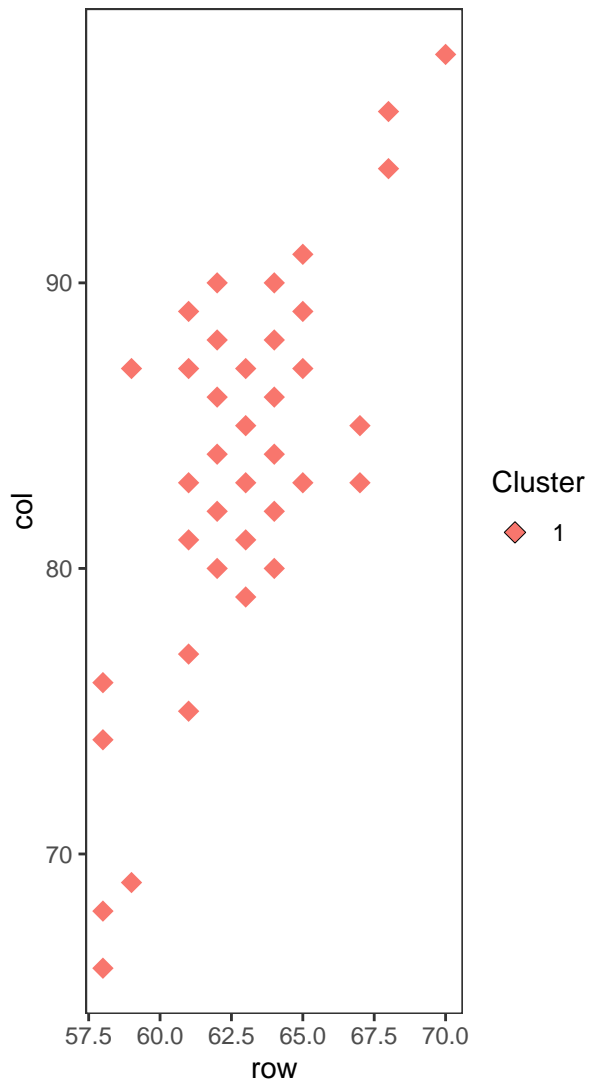
Geneset=Overlap, method=AUCCell



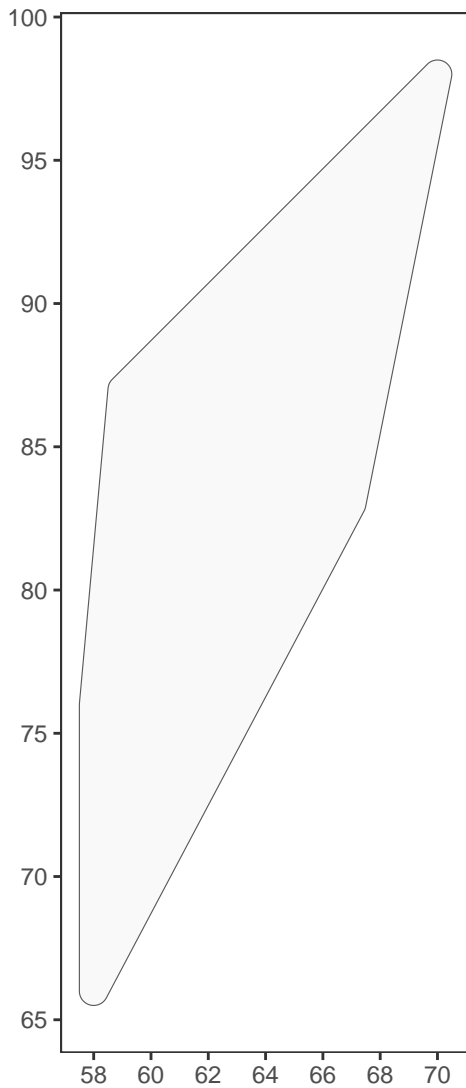
TLS cutoff is 0.257



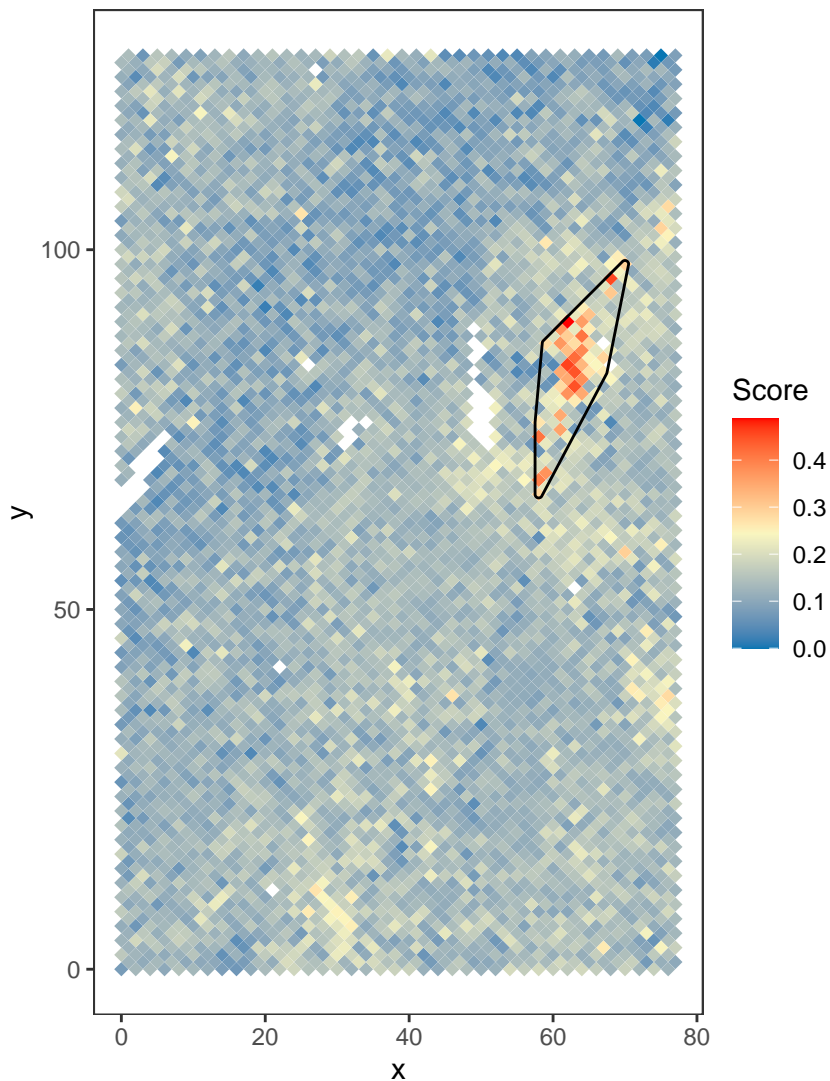
Cluster for TLS prediction



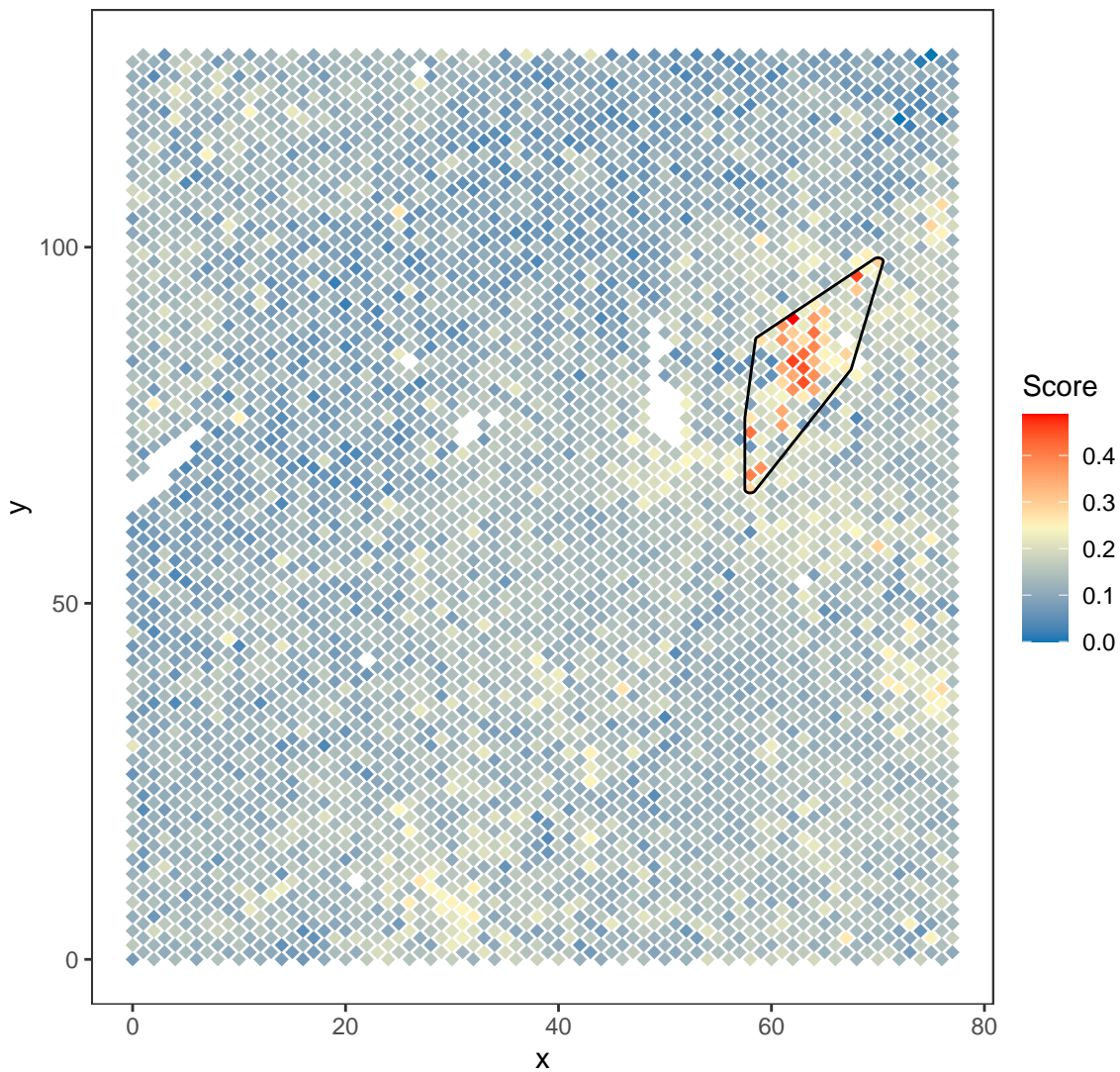
TLS area =1:193.4



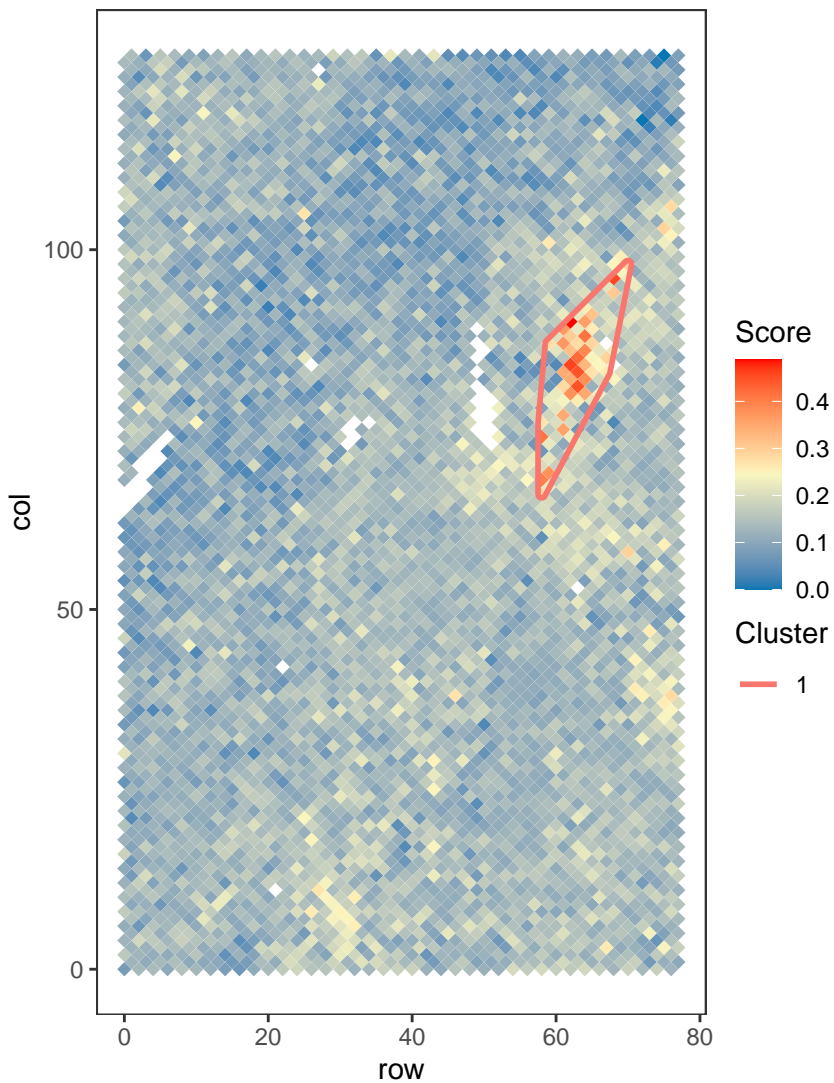
TLS candidate region

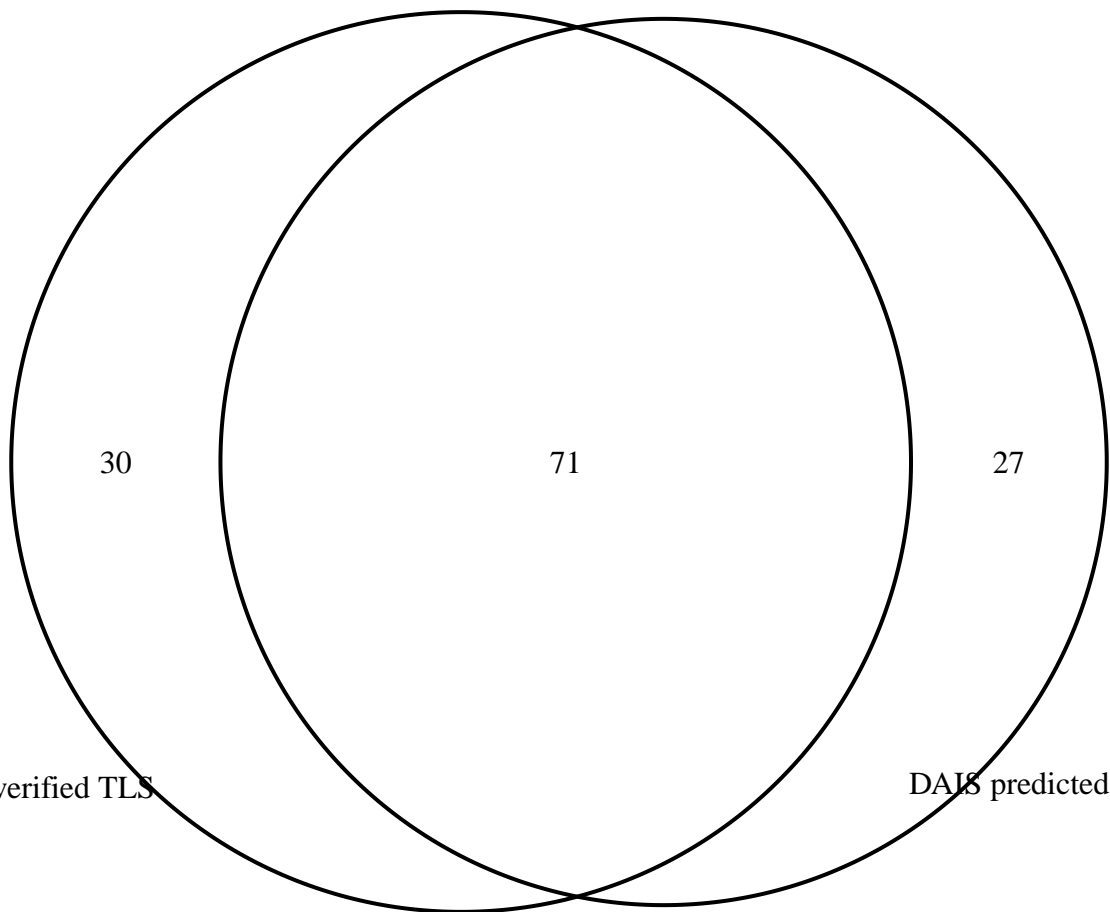


TLS candidate region



k=4,eps=6





verified TLS

DAIS predicted species