

AGCT Tutorial

Step-by-step introduction

AGCT dataset

Species Yeast.txt											
E (entity)	1	Cell1									
EG (entity grouping)	1	1									
T (time)	36	1	2	3	4	5	6	7	8	9	10
R (replicate)	1										11
Data	564										
10001_at	1.994324565	1.805780292	1.173666358	1.147356629	0.940516353	0.930283248					
10003_f_at	22.51645851	11.07745647	19.84449577	17.49943733	17.20314407	18.38671112					
10005_at	2.220204353	1.906358242	2.366628885	1.8436445	1.734006763	1.860566497					
10006_at	0.657207727	0.635838151	0.656072617	0.905511856	1.127946138	1.001089334					
10010_at	29.91146469	26.66589737	32.16685486	22.47244263	15.67227936	10.69390011					
10011_at	0.264472187	0.116763011	0.17026107	0.15185602	0.14927049	0.077342048					
10012_at	1.585698009	3.213872671	3.719636917	3.665916681	4.057239056	4.177559853					
10013_at	0.700340509	0.897109807	1.055618644	0.942632198	0.940516353	1	0.97				
10015_at	4.947786808	9.030056953	14.13961411	9.087739944	9.347924232	6.636165619					
10016_at	3.312145233	2.530635595	2.947786808	3.011248827	2.66554451	2.198257208					
10018_at	43.29738998	25.93988419	41.94778824	43.53430939	36.99775696	33.83224487					
10019_at	40.52213287	39.93179321	34.62996674	26.75140572	24.3254776	23.30827904					
10021_at	0.60839951	0.893641651	0.679909229	0.782902122	0.912458003	0.751634002					
10026_at	1.220204353	2.546820879	1.316685557	2.213723421	2.059483767	1.745097995					
10027_at	6.491487503	10.24624252	7.511918068	8.645668983	8.375982285	7.60566473					
10028_at	0.648127139	0.582658947	0.740068078	0.373453349	0.718294084	0.698257089					
10029_at	2.457434654	2.38959527	2.039727449	1.480315089	0.854096532	0.957516372					
10031_at	1.404086232	1.365317822	1.769580007	1.519685149	1.904601574	1.624183059					
10032_at	0.936435878	1.253179193	1.657207727	1.600674987	1.722783446	1.318082809					
10036_at	1.644721866	1.964161754	2.013620853	2.424072027	2.099887848	2.165577412					
10040_at	1.405221343	1.531791925	1.38138473	1.394825697	1.564534187	1.310457587					
10043_at	9.01816082	19.66589546	13.46765041	11.17660332	11.27833939	9.603485107					
10044_at	2.707150936	1.81734097	1.460839987	1.646794081	1.404040456	1.36601305					
10045_at	2.922815084	8.425433159	45.61407471	69.77278137	60.65769196	49.42483902					
10046_at	0.124858119	0.171098262	0.246311024	0.269966245	0.407407403	0.352941185					
10048_at	1.293984175	1.376878619	1.248581171	1.933633208	1.758698106	2.035947561					
10049_at	1.916004539	1.108670473	0.555051088	0.73453325	0.59932667	0.636165619					
10050_at	3.827468872	4.931791782	3.595913649	5.411698818	2.720538855	5.583877563					
10051_at	1.748013616	2.60346818	1.782065868	2.811023712	2.499439001	2.824618578					
10052_at	1.860385895	1.643930554	1.703745842	1.830146194	1.493827343	1.503268003					
10053_at	8.368899345	4.337572098	9.069239616	7.421822071	9.430976868	6.896513939					
10054_at	23.40522194	23.17687798	29.45857048	19.60517502	17.40067482	11.58060932					
10055_at	8.177071571	4.02543354	2.634506226	2.452193499	2.268238068	1.979302764					
10056_at	2.304199696	5.123699665	3.113507271	4.482564926	4.317620754	3.967320442					
10058_at	2.39160037	5.872832298	6.331441402	7.707536697	8.800224304	8.18954277					
10059_at	4.858115673	4.272832394	3.374574184	4.865016937	5.373737335	4.647058964					
10060_at	3.417707205	12.13410378	9.771850586	11.81439877	9.334456444	8.826797485					
10061_at	42.83314514	49.29364014	31.3689003	38.09674072	32.96408844	31.83660316					
10068_i_at	14.54370117	7.721387386	13.72077274	16.03599548	18.19079781	15.3932457					
10069_f_at	9.527810097	6.567629814	11.72304249	11.0731163	10.35241413	8.009803772					
10071_at	0.402951181	0.436994195	0.712826312	0.512935877	0.586980939	0.412854046					
10072_at	0.473325789	1.161849737	1.089670897	1.17660296	1.125701547	1.058823466					

Species - specify Gene Ontology File

E (entity) - number of sets (cells here) to analyze

EG (entity grouping) - number of subgroups (optional, can be ignored)

T (time) - total number of timestamps, followed by a list of stamps

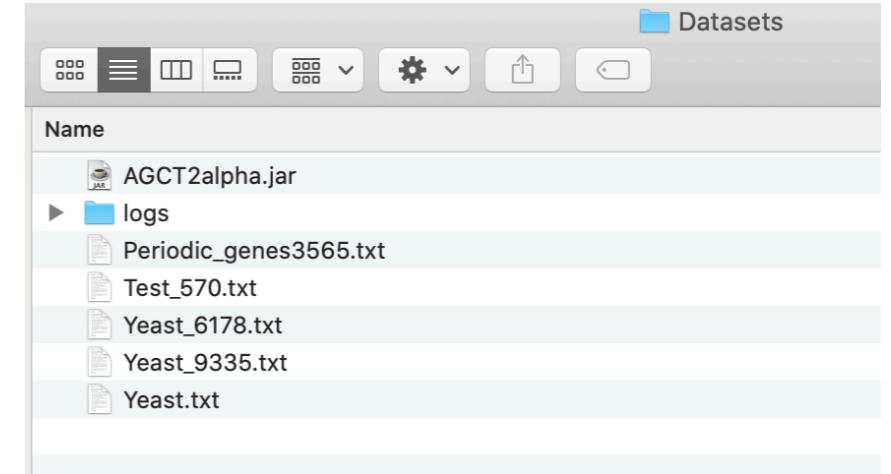
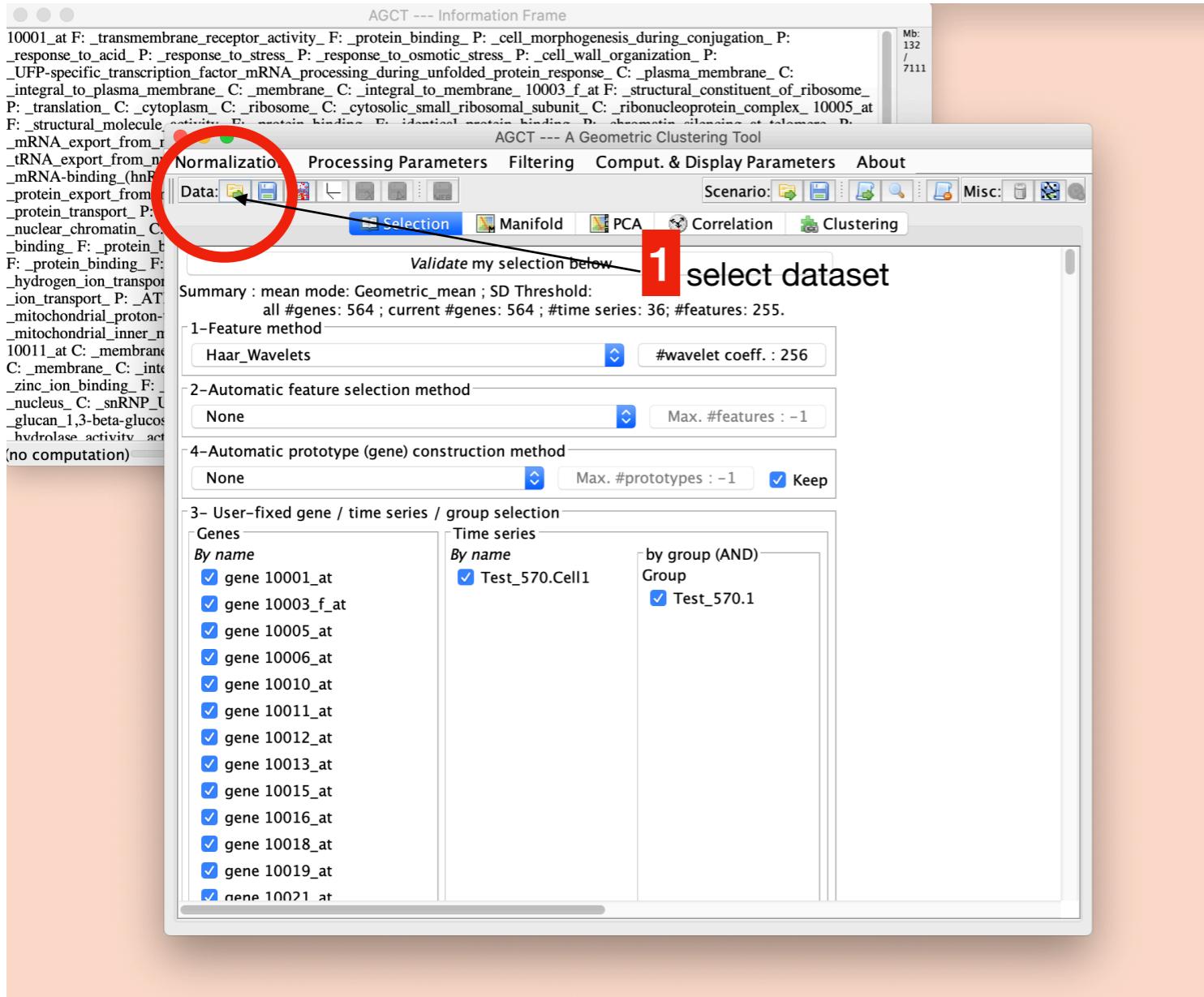
R (replicate) - number of replicates

Data - number of entities in the list (number of probes here)

List of IDs followed by values (in this case: E*EG*T*R=36 tags)

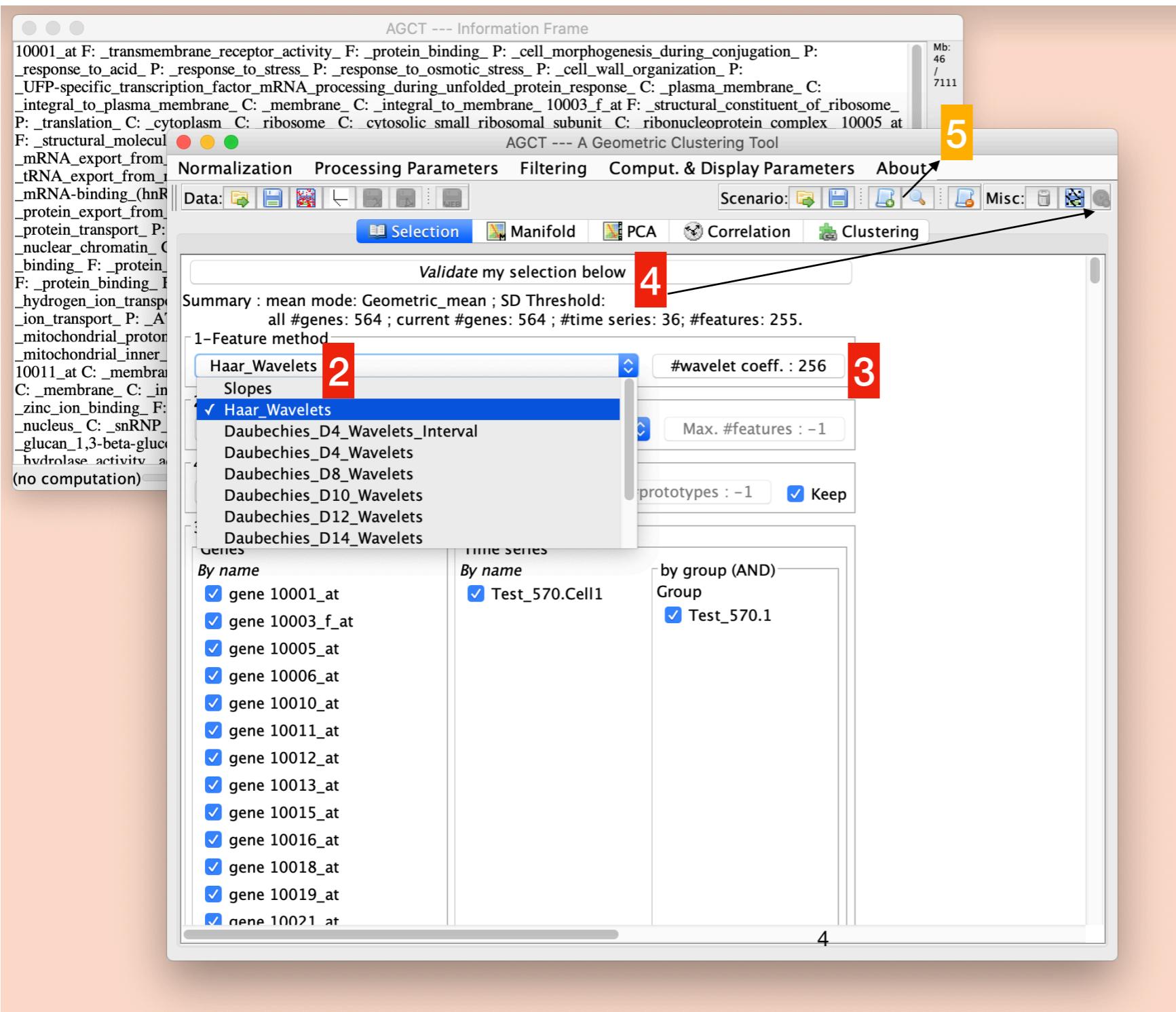
10001_at 1.99432457 1.80578029 1.17366636 1.14735663 ...n=36

Launch AGCT and loading a dataset

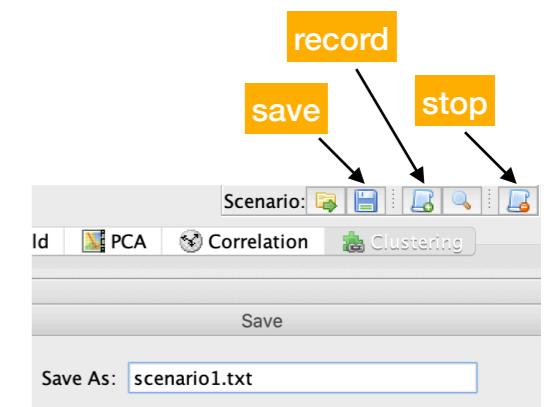


```
java -d64 -Xmx8000m -jar AGCT2alpha.jar
```

Execute AGCT on dataset

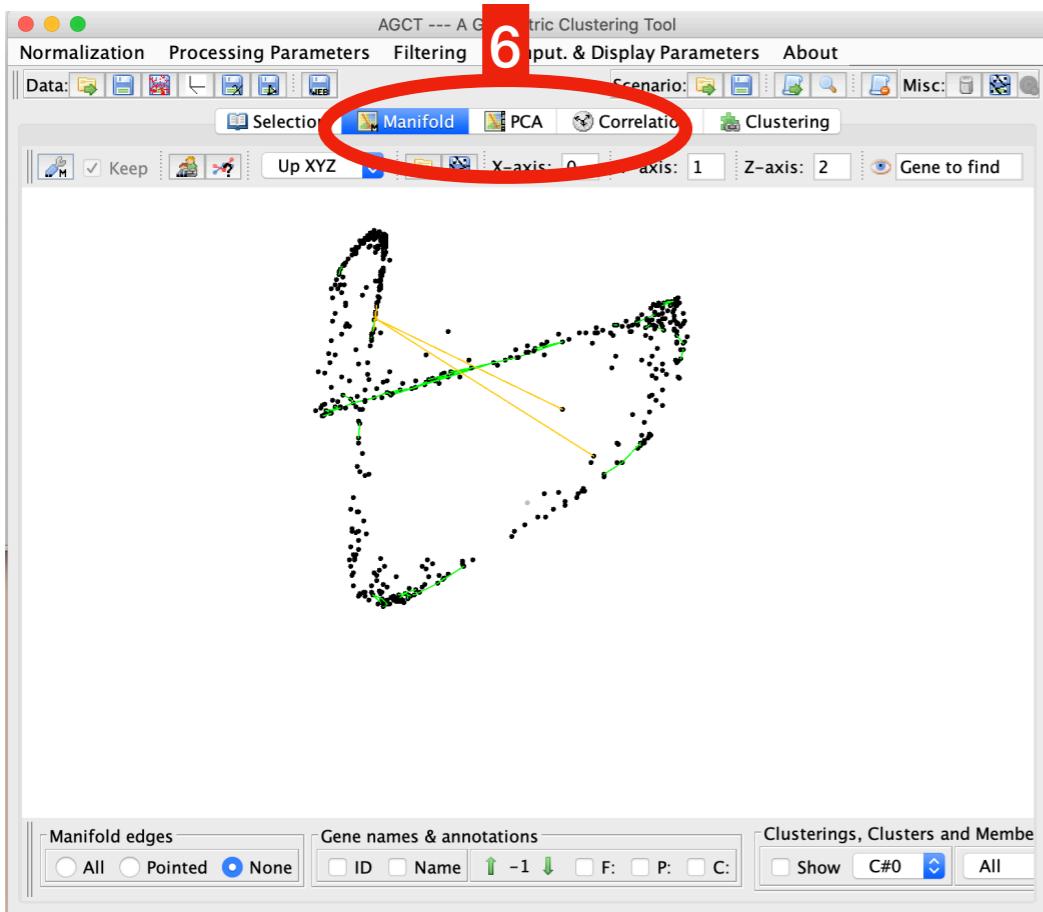


- 2 Select feature compression method
- 3 Select wavelet coefficient (if used)
- 4 Execute job
- 5 Scenario automatically recorded before manually stopped

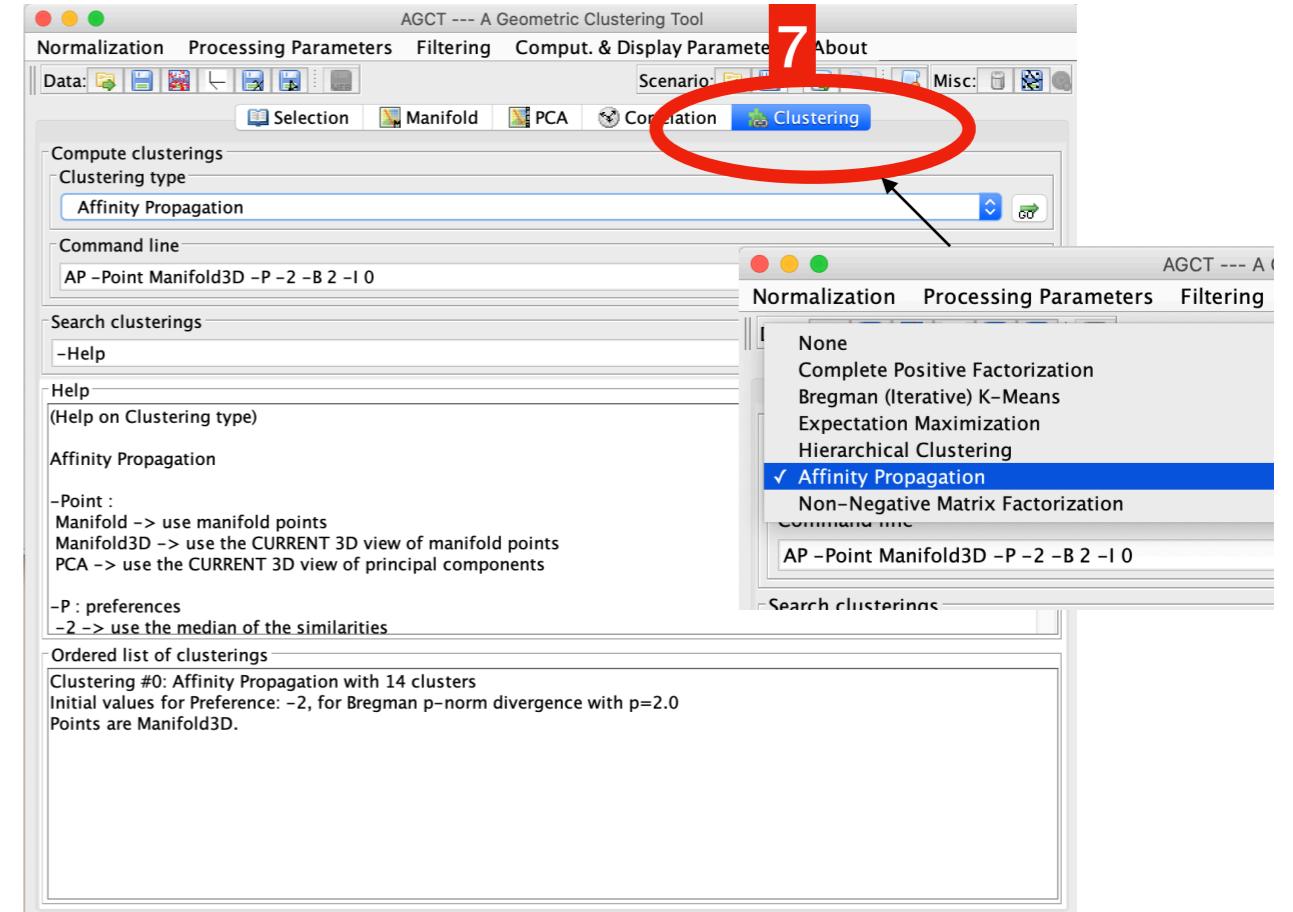


The scenario can be saved to manifold calculation.
Next time, to load **your recorded scenario** you have, first, **to stop the automatic recording** scenario button.

Visualization Manifold/PCA



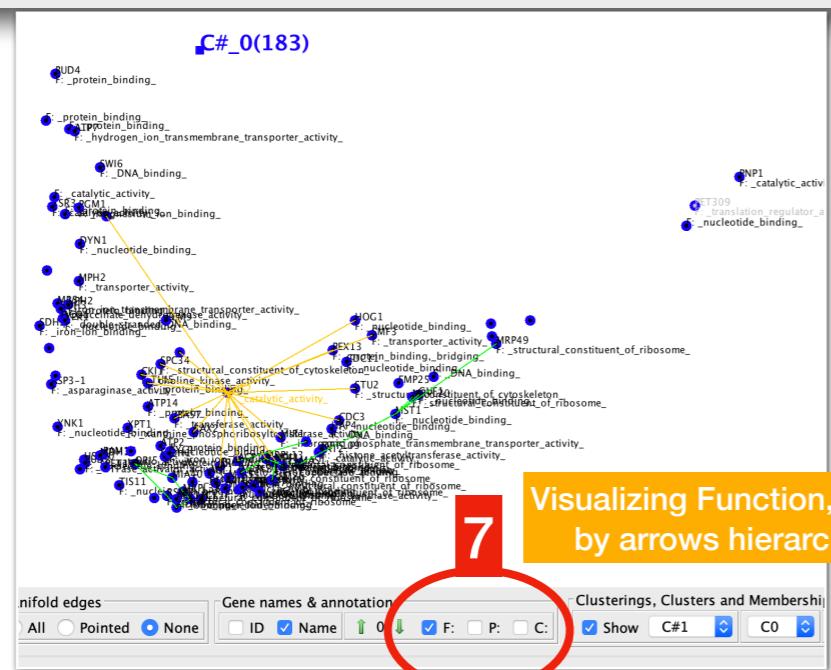
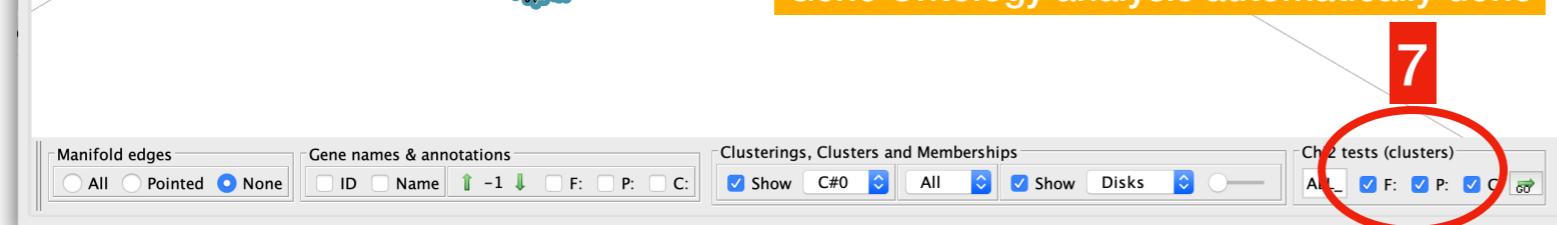
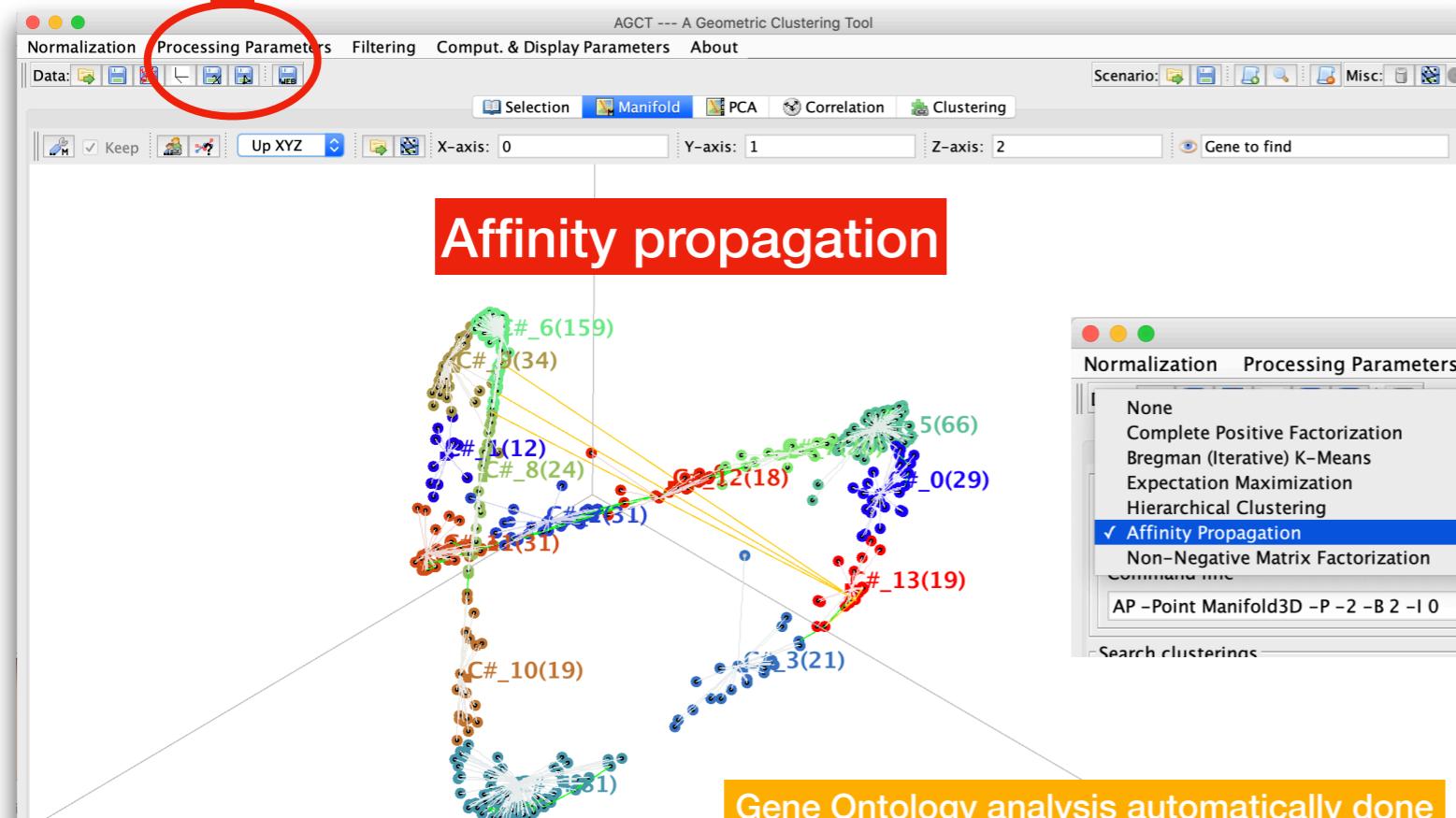
Clustering method selection



Gene Ontology analysis saved (.txt)

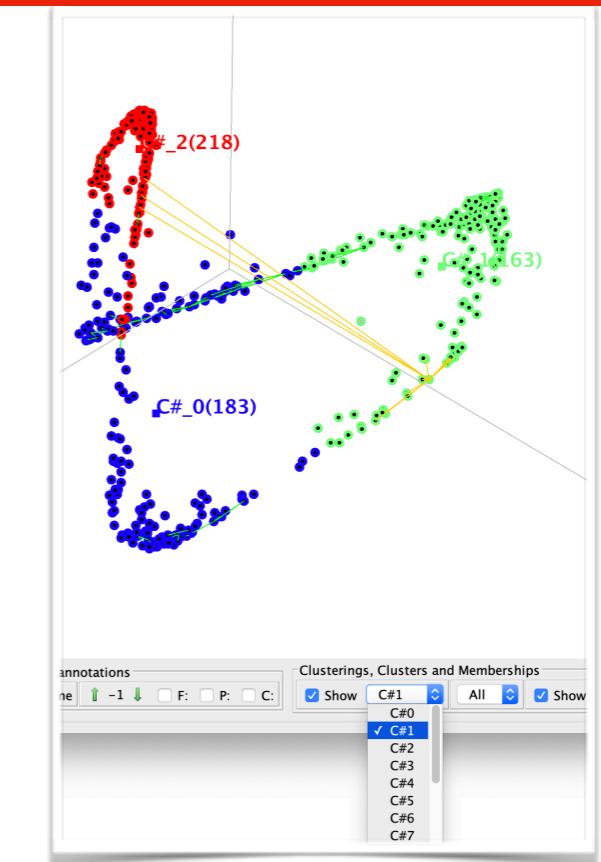
Results

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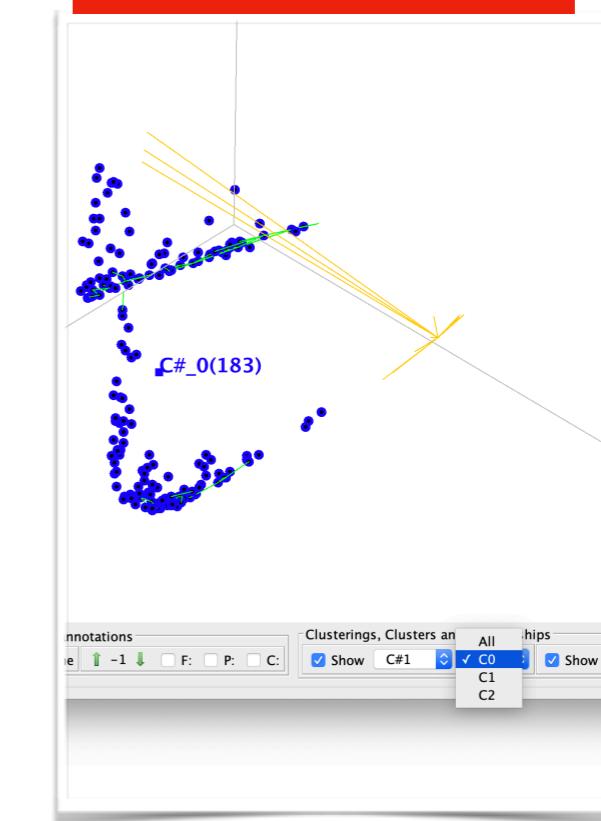


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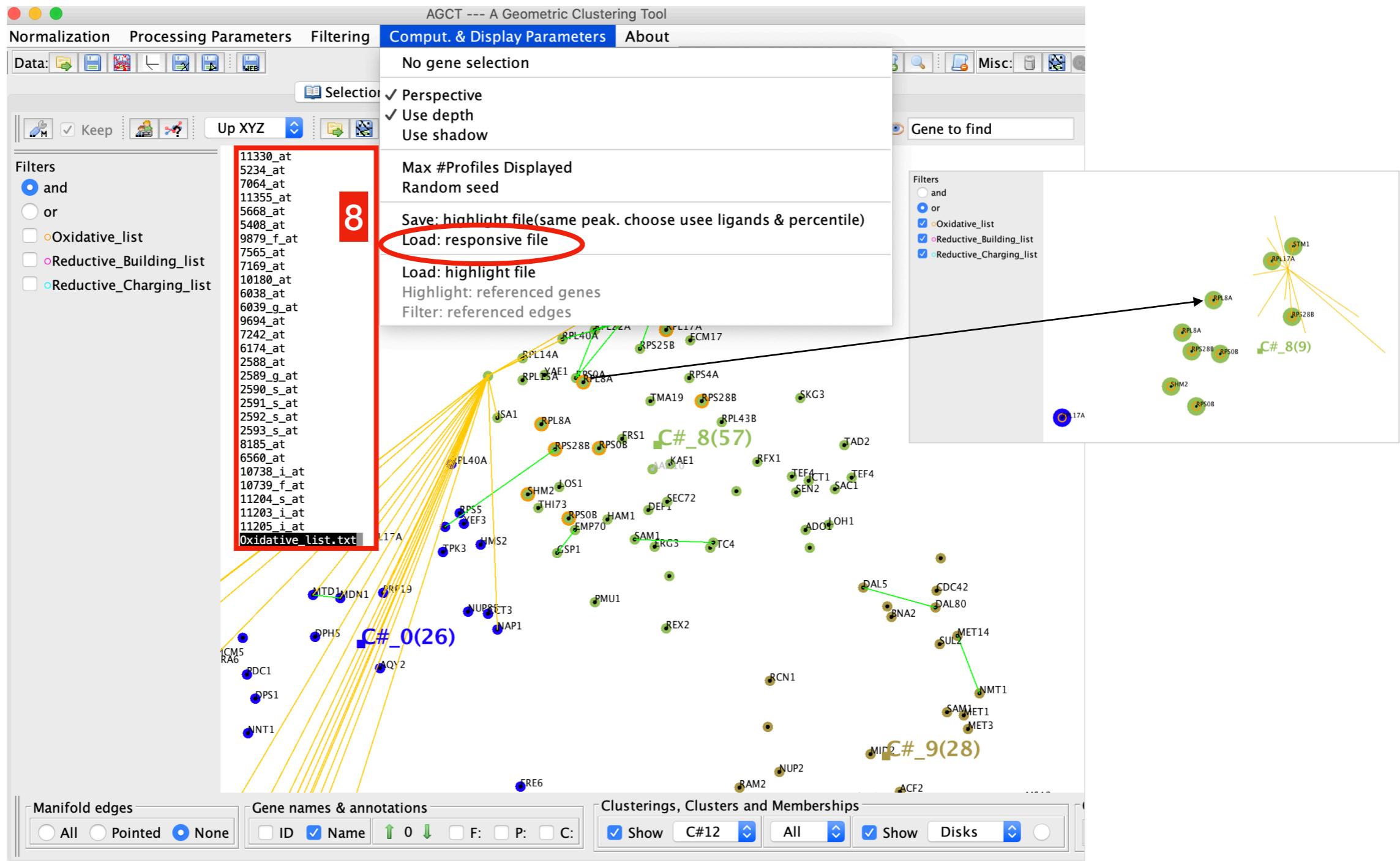
Bregman K-means (k=3)



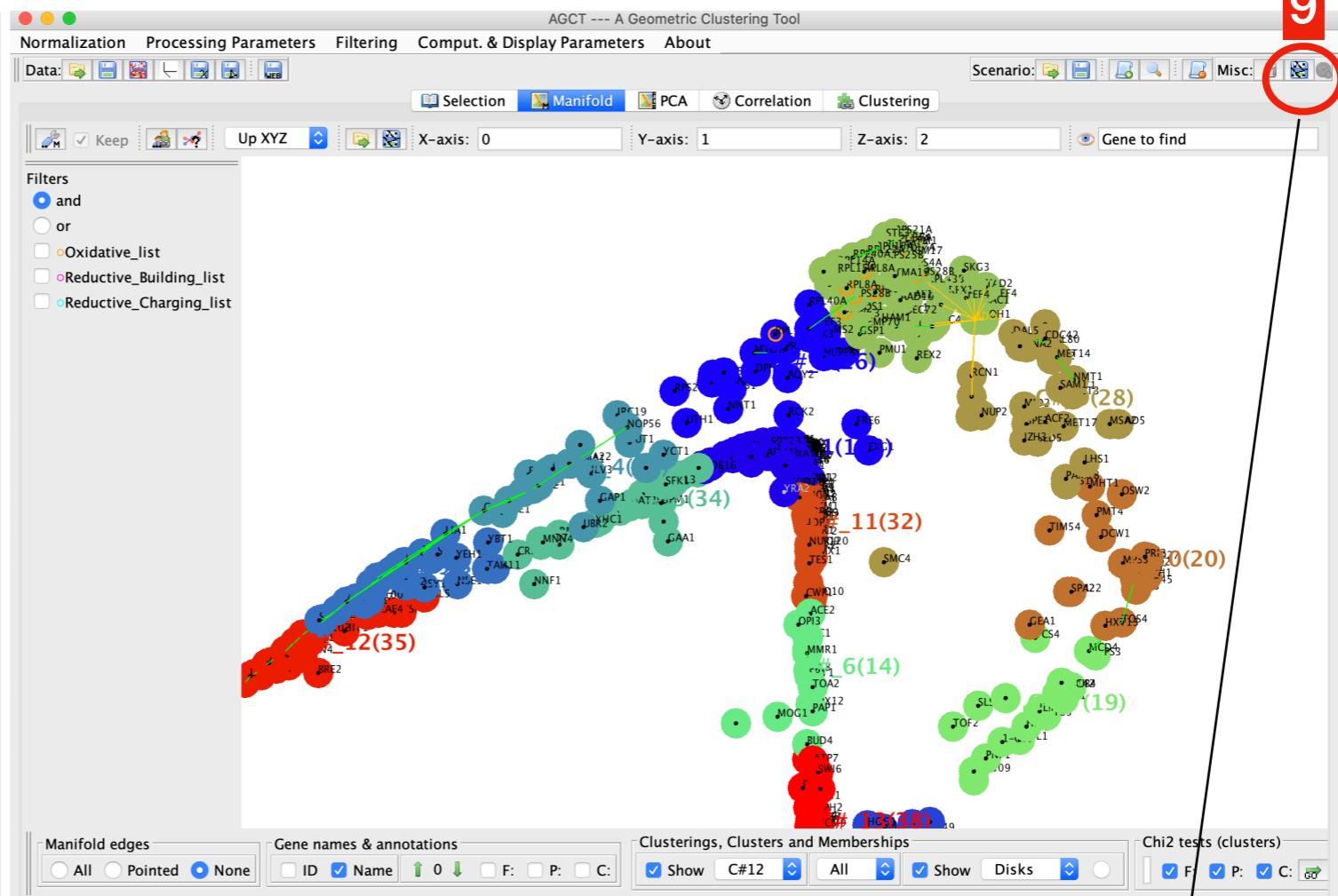
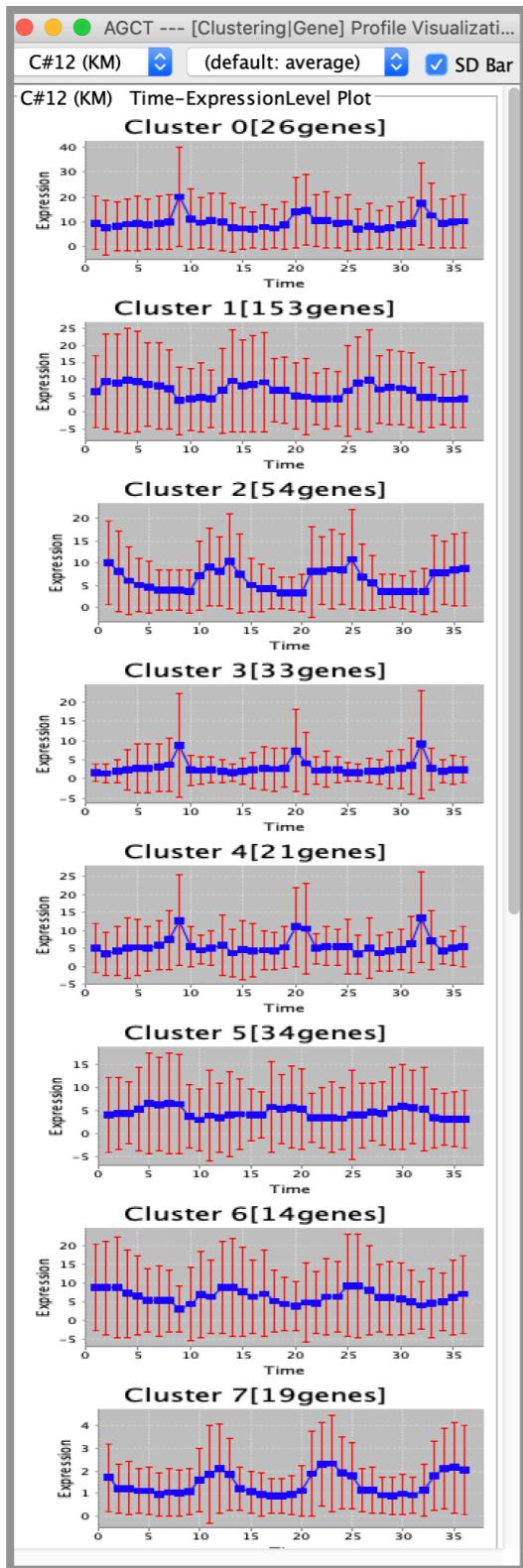
Select one cluster



Loading genes to highlight from separate file



Cluster profiles



Cluster saving

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AGCT2alpha.jar	Test_570_cluster_4.txt
Periodic_genes3565.txt	Test_570_cluster_5.txt
Test_570.txt	Test_570_cluster_6.txt
Test_570_clusterProfile.txt	Test_570_cluster_7.txt
Test_570_cluster_1.txt	Test_570_cluster_8.txt
Test_570_cluster_10.txt	Test_570_cluster_9.txt
Test_570_cluster_11.txt	Test_570_grouping.txt
Test_570_cluster_12.txt	Yeast.txt
Test_570_cluster_13.txt	Yeast_6178.txt
Test_570_cluster_14.txt	Yeast_9335.txt
Test_570_cluster_2.txt	logs
Test_570_cluster_3.txt	