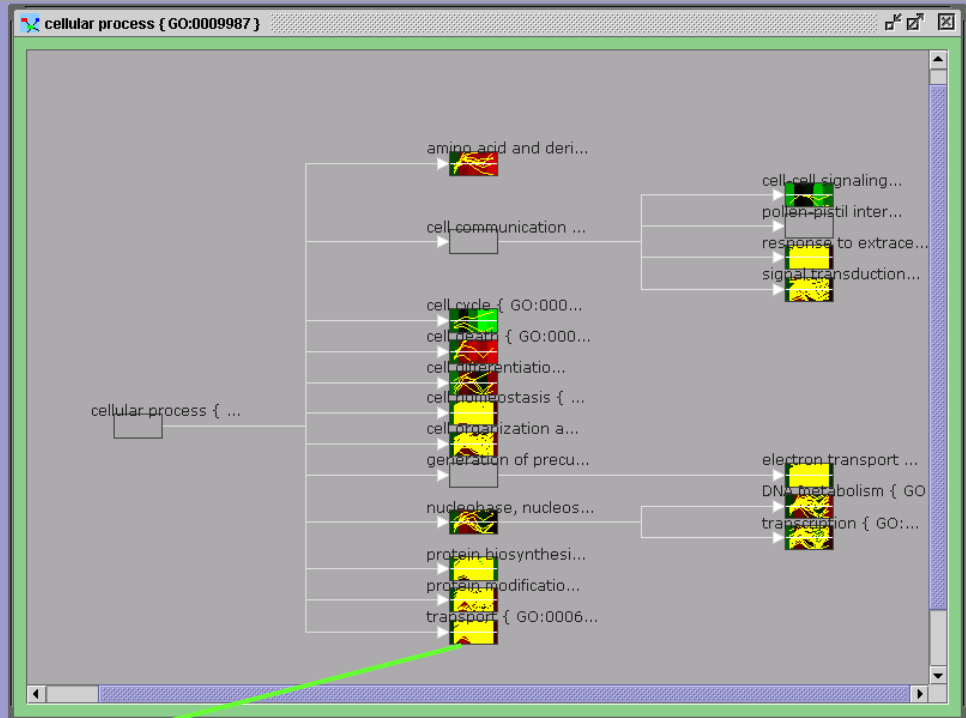
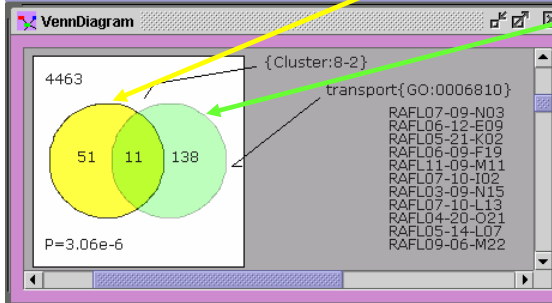
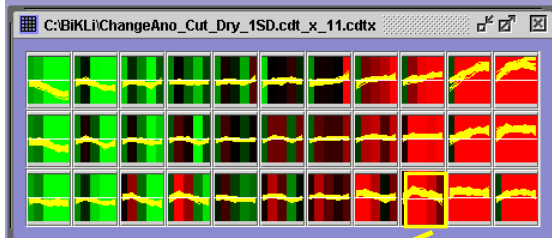


C:\BIKLiChangeAno_Cut_Dry_1SD.cdt_x_11.cdtx

- biological_process(GO_SUBSET0008150)
 - DRY_1
 - DRY_2
 - DRY_5
 - DRY_10
 - DRY_24
- biological_process(GO:0008150)
 - behavior(GO:0007610)
 - biological_process unknown(GO:0000004)
 - cellular_process(GO:0009987)
 - development(GO:0007275)
 - physiological process(GO:0007582)
- cellular_component(GO_SUBSET0005575)
- molecular_function(GO_SUBSET0003674)



Result

Target	Query	A	B	C	D	Phi	P	#	P'
endosome GO:0005788	Cluster:8-2	19	43	366	4235	0.09	2.61e-7	1	2.61e-7
transport GO:0006810	Cluster:8-2	11	51	138	4463	0.09	3.06e-6	1	3.06e-6
death GO:0016265	Cluster:8-2	5	57	97	4504	0.04	0.01	1	0.01
cellular_component GO:0005575	Cluster:8-2	40	22	2250	2351	0.03	0.01	1	0.01
Golgi apparatus GO:0005794	Cluster:8-2	3	59	44	4557	0.04	0.02	1	0.02
protein binding GO:0005515,GO:0045308	Cluster:8-2	4	58	85	4516	0.03	0.02	1	0.02
cytoplasm GO:0005737	Cluster:8-2	27	35	1417	3184	0.03	0.03	1	0.03
cell GO:0005623	Cluster:8-2	35	27	1986	2615	0.03	0.03	1	0.03
protein modification GO:0006464	Cluster:8-2	3	59	58	4543	0.03	0.04	1	0.04