



• MEV软件无需安装,无需解压,但是需要在使用该软件之前安装JAVA,点击文件夹中的JAVA软件 Java Runtime Environment-6.0.330.exe 安装即可。

1 打开MeV,导入数据。



导入文件格式.txt

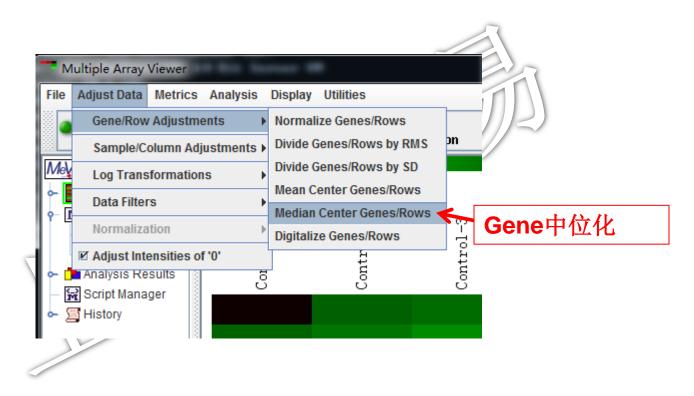
基因或miRNA名称, 可以是probe ID,gene symbol,miRNA等

相应样本的标准化 信号值

A	В	С	D	E	F	G	Н	I	J	K
<u>systemat:</u>	Control-1	Control-2	Control-3	Control-4	Control-5	Treatment	Treatment	Treatment	Treatment	Treatmen
hsa-miR-	4.117472	2.425547	2.257792	2.721742	2.507492	5.290645	3.963713	4.381024	3.795941	6.483124
hsa-miR-	2.392147	2.093825	1.76745	2.318852	2.215987	4.488134	3.37224	2.601644	3.026403	3.770224
hsa-miR-:	0.910198	1.103413	0.961108	0.312955	0.513069	1.673431	1.927288	1.155438	1.118214	3.129052
hsa-miR-:	-5.67499	-0.84145	-6.22027	-6.38511	-1.04084	2.513427	1.046393	0.565252	2.219901	2.718564
hsa-miR-:	1.711456	2.342337	2.020065	0.669011	1.49876	4.628832	2.713693	4.554214	1.211871	3.902363
hsa-miR-:	10.54308	10.47967	10.6221	9.991513	10.17862	12.3105	11.5511	12.04947	11.66502	12.63161
hsa-miR-:	1.884155	3.448479	2.421772	1.723666	2.070822	4.666307	4.214134	6.09797	3.218871	4.289248
hsa-miR-:	-5.67499	-1.66584	-6.22027	-2.06629	-2.19858	-1.35716	0.121457	0.158635	-1.13201	-1.47737
hsa-miR-	-0.834	-0.07796	-0.17079	-0.32886	-1.05529	0.772286	1.249588	1.166869	-0.56126	0.160276
hsa-miR-	-5.67499	-6.31927	-6.22027	-6.38511	-5.99462	-1.25885	-0.31737	-0.90772	-1.07143	-0.87847
hsa-miR-	-5.67499	-6.31927	-0.15489	-6.38511	-0.99414	-0.45932	-0.0796	1.881273	-1.3304	0.409467
hsa-miR-	3.472245	2.72126	3.170607	3.223963	3.225354	4.588028	3.088008	5.023236	4.168779	6.334113
hsa-miR-	-5.67499	-6.31927	-1.05804	-1.50603	-5.99462	0.769371	-0.19699	0.127119	-1.0875	-0.63399
hsa-miR-	-5.67499	-1.14248	-6.22027	-6.38511	-5.99462	1.936681	0.538888	0.359389	0.004417	0.062908
hsa-miR-	-5.67499	-0.88969	-1.1131	-6.38511	-5.99462	0.800286	1.214684	-0.07767	-1.6754	1.259772
hsa-miR-	0.488954	1.203315	0.091881	0.428859	0.843842	3.048372	2.580061	4.394995	0.310273	3.270484
hsa-miR-	1.795681	1.449729	1.606758	1.492504	1.749229	3.52921	3.385056	2.425678	1.471901	2.964652
hsa-miR-	1.876844	1.276551	1.274255	1.323378	1.066724	4.5613	3.337945	1.182763	3.031988	3.088815
hsv1-miR	-5.67499	-6.31927	-6.22027	-1.4512	-5.99462	-0.82637	0.063505	1.245648	-0.29145	-0.65783
hsa-let-	10.7838	10.10293	10.30817	10.73809	10.68266	9.630089	8.378316	7.681264	9.886524	10.39732
hsa-let-	9.819313	9.510312	9.415765	10.09319	9.918474	8.559124	7.143523	8.140289	8.901421	9.530088
hsa-let-	8.800498	8.444374	8.495061	8.922276	8.715842	7.154	5.709683	6.616036	7.707176	7.788385
hsa-let-	6.920078	6.124488	6.245704	7.097914	6.937068	5.180437	4.221624	4.135463	6.516961	5.744849
hsa-miR-:	7.348121	5.966193	6.464092	6.120363	6.996758	3.216342	1.678957	-6.18626	4.55841	2.656587
hsa-miR-	7.038822	6. 478651	6.640234	7.128747	7.165916	5.047347	2.804562	3.45001	5.40336	4.579755



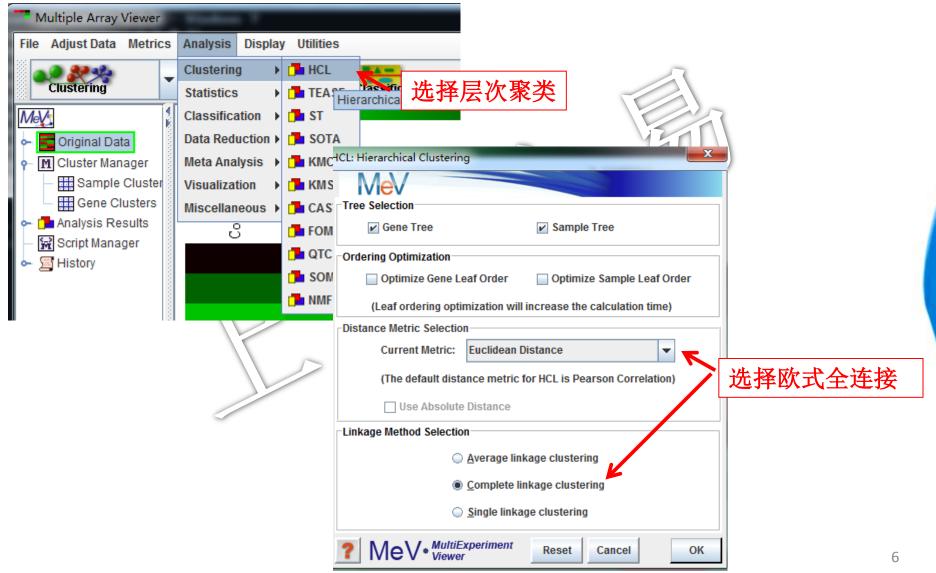
2 数据中位化(一般需对gene进行中位化)

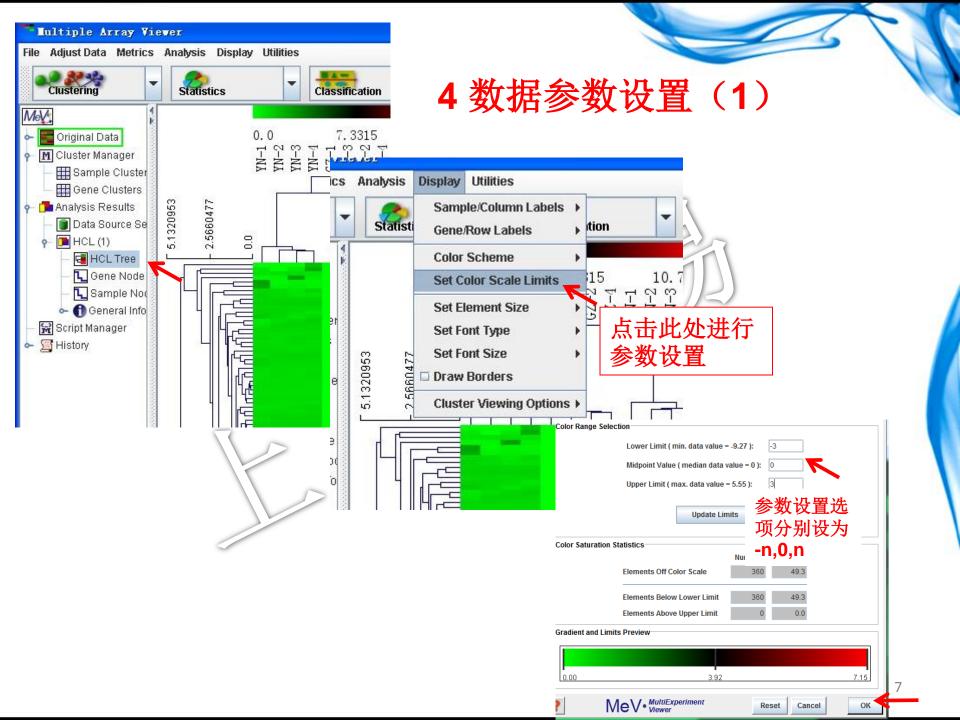


注意: 该步骤点击完成软件中不会出现任何变化, 但实际上内部已经运 算完成,切忌重复点击。

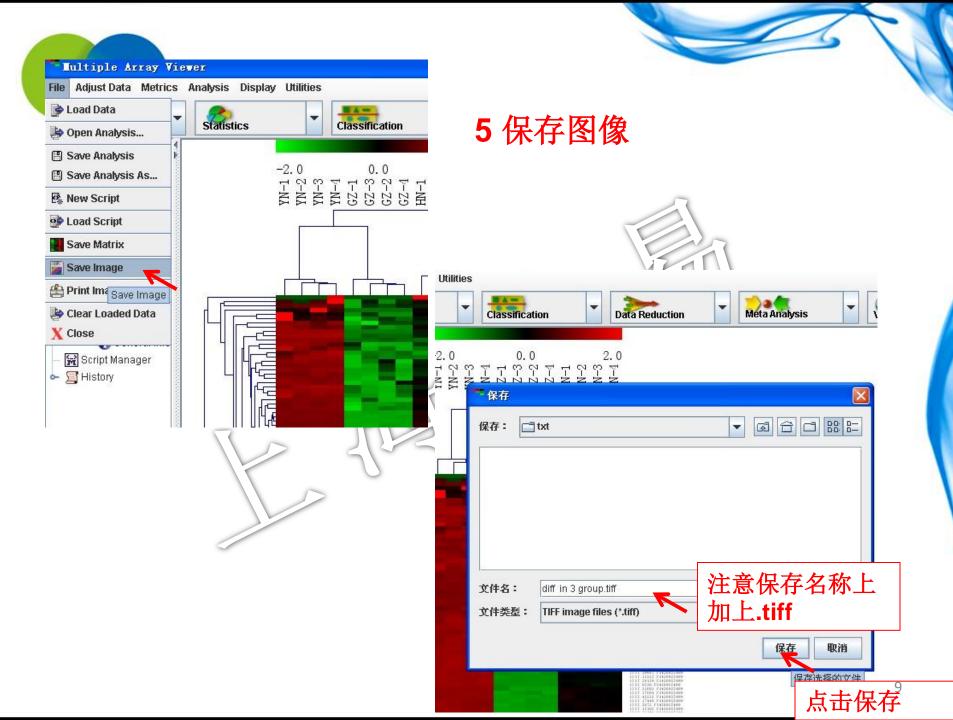


3 进行聚类分析



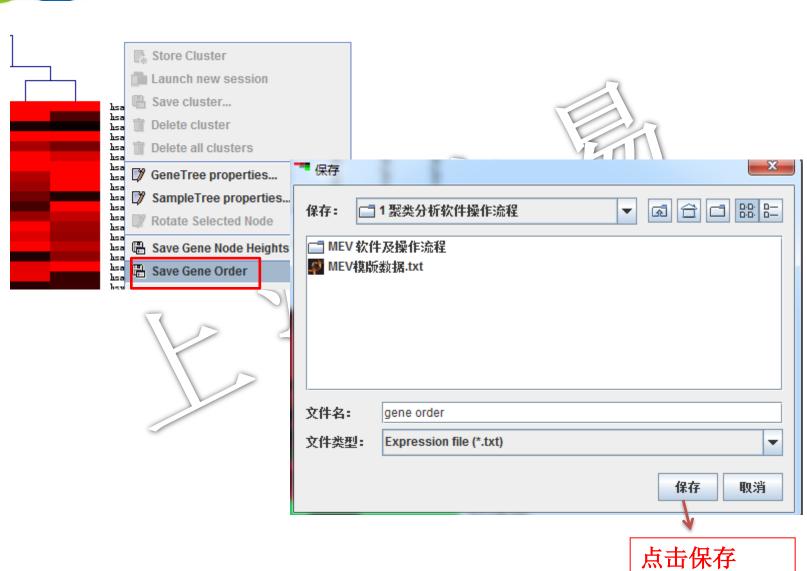








6. 保存gene order





聚类后的位置情况

原来的位置

