

DNA Microarrays

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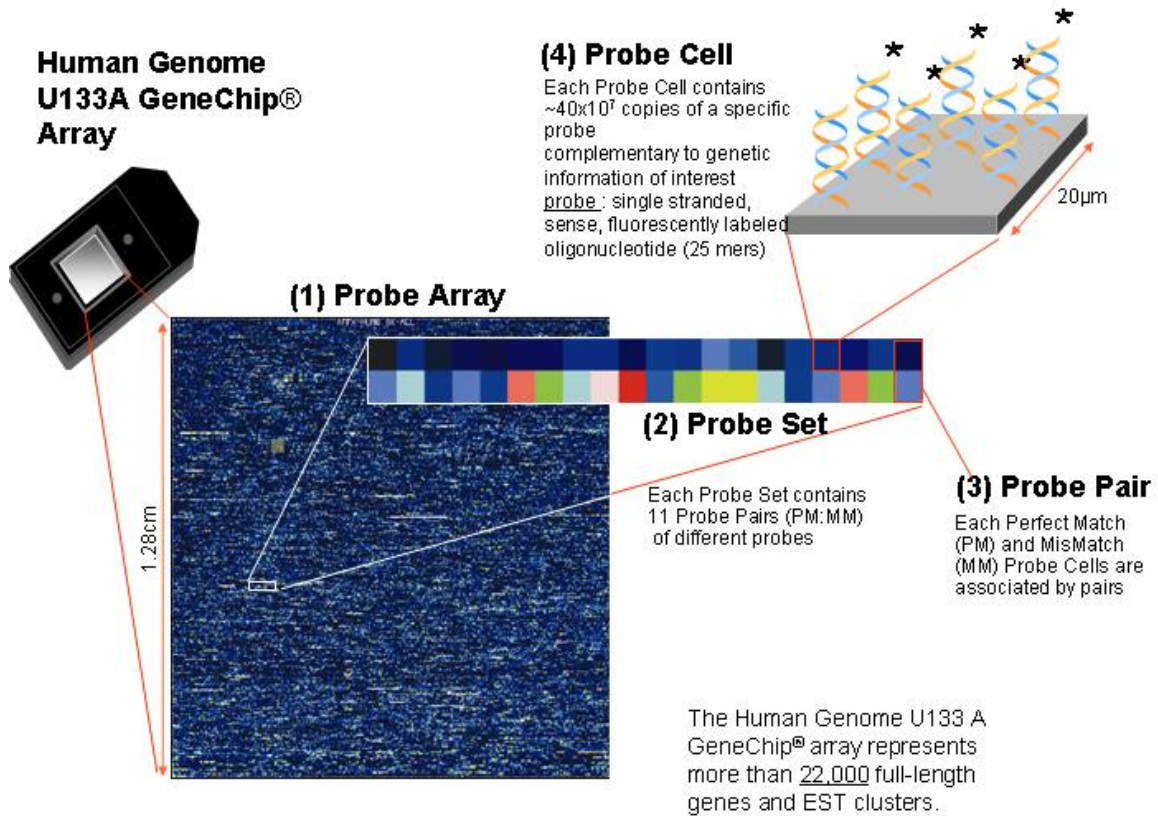
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Microarrays

DNA Microarrays



Un ejemplo

- [gse21779](#)

ExpressionSet

```
pacman::p_load(ALL)
```

```
data(ALL)
```

La clase del objeto la podemos ver con

```
class(ALL)
```

```
[1] "ExpressionSet"  
attr(,"package")  
[1] "Biobase"
```

El método `print` nos da un breve resumen de la clase.

```
ALL
```

La matriz de expresión la obtenemos con el accessor `Biobase::exprs`.

```
      01005      01010      03002      04006      04007      04008      04010  
1000_at  7.597323  7.479445  7.567593  7.384684  7.905312  7.065914  7.474537  
1001_at  5.046194  4.932537  4.799294  4.922627  4.844565  5.147762  5.122518  
1002_f_at 3.900466  4.208155  3.886169  4.206798  3.416923  3.945869  4.150506  
1003_s_at 5.903856  6.169024  5.860459  6.116890  5.687997  6.208061  6.292713  
1004_at  5.925260  5.912780  5.893209  6.170245  5.615210  5.923487  6.046607  
1005_at  8.570990 10.428299  9.616713  9.937155  9.983809 10.063484 10.662059  
      04016      06002      08001      08011      08012      08018      08024  
1000_at  7.536119  7.183331  7.735545  7.591498  7.824284  7.231814  7.879988  
1001_at  5.016132  5.288943  4.633217  4.583148  4.685951  5.059300  4.830464  
1002_f_at 3.576360  3.900935  3.630190  3.609112  3.902139  3.804705  3.862914  
1003_s_at 5.665991  5.842326  5.875375  5.733157  5.762857  5.770914  6.079410  
1004_at  5.738218  5.994515  5.748350  5.922568  5.679899  6.044520  6.057632  
1005_at 11.269115  8.812869 10.165159  9.381072  8.227970  7.627248  7.667445  
      09008      09017      11005      12006      12007      12012      12019  
1000_at  7.891793  7.756734  7.640012  7.759599  7.678636  7.464285  7.652719  
1001_at  5.999496  4.987595  4.967288  4.770481  5.456332  4.785863  5.175609  
1002_f_at 4.001606  4.048901  3.796550  3.912707  3.870893  3.930832  3.932360  
1003_s_at 5.832952  6.097900  6.094379  6.235795  5.971466  6.037364  6.194623  
1004_at  5.717497  6.210092  5.751805  5.883340  5.918456  5.725421  5.969027  
1005_at 10.206353 10.015466  9.358516  8.824348  9.262478  7.232927 10.243610  
      12026      14016      15001      15004      15005      16004      16009  
1000_at  7.501591  7.570417  7.331509  7.366208  7.455451  7.328875  7.297313  
1001_at  5.188992  5.258312  4.627955  4.733495  5.125098  5.332775  5.215707  
1002_f_at 4.188444  4.028859  4.099497  3.831617  3.786741  4.648154  4.176687  
1003_s_at 6.231228  6.348593  6.057790  6.043205  6.303592  5.905220  6.274415  
1004_at  6.357476  6.173530  5.729543  5.671164  5.977084  5.721572  6.000902  
1005_at  7.808452  7.557919 10.233185  9.749860  8.477219  9.884608  9.599149
```

	19005	20002	22009	22010	22011	22013	24001
1000_at	7.563561	7.541133	8.016818	7.862181	7.702580	7.412003	7.916169
1001_at	4.858392	4.964424	5.216252	5.135825	4.802946	5.222676	4.790170
1002_f_at	4.097332	3.789888	3.980839	3.954917	3.971934	4.109899	3.899038
1003_s_at	5.838829	6.160238	6.343042	6.195307	5.865581	6.243157	6.022905
1004_at	5.970021	6.054232	6.024327	6.114502	6.035582	5.896131	5.800271
1005_at	9.683996	9.221405	8.571812	8.841628	8.489550	8.998592	8.933302
	24005	24008	24010	24011	24017	24018	24019
1000_at	7.595848	7.296349	7.506236	7.144425	7.513972	7.815971	7.406135
1001_at	4.804743	5.002518	4.218220	5.228892	5.264158	4.899316	4.791335
1002_f_at	3.677240	3.906343	3.579385	3.829513	3.965467	4.058576	4.131234
1003_s_at	5.669771	5.668509	5.273965	5.817272	6.088179	6.387995	6.343673
1004_at	5.475892	5.437155	4.634124	5.552223	5.982065	5.874817	6.048209
1005_at	9.626267	10.826157	8.782330	7.881855	11.069535	9.102971	10.037809
	24022	25003	25006	26001	26003	26005	26008
1000_at	7.300980	7.845054	7.651229	7.376930	7.663977	7.250353	7.663612
1001_at	5.177703	5.250315	4.896195	5.123546	5.078104	4.945670	5.124591
1002_f_at	3.838198	4.046442	4.120495	4.131492	3.803233	4.105035	4.119660
1003_s_at	5.863318	6.205917	6.298788	6.118064	6.199316	5.782105	6.271931
1004_at	5.669051	5.931859	5.915944	6.002812	5.822230	5.930979	6.073099
1005_at	9.095296	8.670866	10.496309	9.046483	9.104846	9.756972	9.071812
	27003	27004	28001	28003	28005	28006	28007
1000_at	7.329996	7.360754	7.035203	7.705260	7.551734	7.538601	7.501531
1001_at	5.438098	4.757900	5.005279	5.009705	4.944978	4.511194	4.888814
1002_f_at	3.677207	3.638739	3.800893	4.238111	3.719482	3.788262	3.626149
1003_s_at	5.899308	5.664813	5.732956	6.379139	5.833428	5.362676	5.702537
1004_at	5.718582	5.595820	5.485361	5.881472	5.554058	4.986320	5.743743
1005_at	6.969776	8.867644	7.067019	9.080559	8.245585	7.807180	10.085771
	28019	28021	28023	28024	28028	28031	28032
1000_at	7.116676	7.107979	7.427808	6.549926	7.514761	7.377215	6.973861
1001_at	5.275964	4.865566	5.057619	5.185277	4.788468	4.778381	4.970430
1002_f_at	4.192648	3.979372	3.791415	3.943834	3.924333	3.657005	3.706332
1003_s_at	6.196541	5.804445	5.719376	5.943116	5.719659	5.939648	5.968072
1004_at	5.926093	5.768851	5.478333	5.756534	5.740607	5.770578	5.905965
1005_at	8.097072	8.661098	9.106441	8.804075	8.235771	10.607653	6.760202
	28035	28036	28037	28042	28043	28044	28047
1000_at	7.227516	7.407561	7.158049	7.235291	7.589310	7.988476	7.362458
1001_at	6.408157	5.042222	5.431469	4.686293	4.851805	4.894379	4.843868
1002_f_at	3.995074	3.714084	4.302001	3.677909	3.831514	3.690856	3.646990
1003_s_at	6.272305	5.733332	6.253362	6.098969	6.132159	6.130691	5.628370
1004_at	6.050495	5.651345	6.494545	6.109255	5.867806	5.592139	5.644372
1005_at	8.957027	8.764321	9.102879	7.008132	7.720932	7.861043	6.642728
	30001	31007	31011	33005	36001	36002	37013

1000_at	7.508667	7.147843	7.651676	7.486432	7.759074	7.473427	7.627685
1001_at	5.587029	4.943857	4.741654	4.642628	4.962544	4.953122	5.358236
1002_f_at	3.765444	3.789166	3.790688	3.682768	3.740679	3.688162	4.008891
1003_s_at	6.078532	5.858604	6.251328	5.961910	5.936883	5.642185	6.314849
1004_at	5.935291	5.526191	5.820374	5.810047	5.616831	5.678327	6.044299
1005_at	9.075910	9.588264	9.585180	11.609025	11.653732	8.893931	7.950866
	43001	43004	43007	43012	48001	49006	57001
1000_at	7.577529	7.600206	7.776844	7.585928	7.450666	7.004613	7.195206
1001_at	5.054157	4.879037	4.949908	5.057530	4.960382	4.836905	4.744006
1002_f_at	3.932435	4.028704	3.689141	3.891536	4.061201	3.699625	3.973128
1003_s_at	6.310934	6.086349	5.658127	6.363734	6.099140	5.616555	5.962672
1004_at	5.782177	5.817414	5.621938	5.975024	5.853644	5.704549	5.765632
1005_at	8.569400	10.693175	7.601647	9.377819	10.929231	9.193089	10.691061
	62001	62002	62003	63001	64001	64002	65005
1000_at	7.407351	7.756195	7.913324	7.270997	7.694588	7.583071	7.609538
1001_at	4.930312	5.238937	5.074681	4.513671	4.928159	4.804083	4.715693
1002_f_at	3.734818	3.945514	3.926906	3.639640	3.806746	4.104208	3.453649
1003_s_at	5.730142	6.061704	6.208286	5.519846	5.834263	6.340025	5.584102
1004_at	5.512776	5.956554	6.028228	5.041052	5.912557	6.056120	5.611407
1005_at	9.108345	9.507559	9.693530	7.758929	8.883131	10.320243	7.757368
	68001	68003	84004	LAL5	01003	01007	02020
1000_at	7.324502	7.545120	7.679603	7.604093	7.240252	7.676749	7.934247
1001_at	5.379102	4.650231	4.795495	4.988922	5.224752	5.129002	5.667907
1002_f_at	4.066075	3.626514	3.554142	3.960894	3.862458	4.169622	4.173444
1003_s_at	6.121059	6.347044	5.471594	5.761373	6.430906	6.202916	6.083160
1004_at	6.224473	5.884682	5.505538	6.048169	5.940335	5.979690	6.127251
1005_at	11.165801	8.986872	9.984865	9.741136	9.790736	8.671713	8.308241
	04018	09002	10005	11002	12008	15006	16002
1000_at	7.874448	7.404271	7.775253	7.771891	7.355677	7.388882	7.589734
1001_at	5.005420	5.127949	4.423445	4.476761	5.461252	5.330129	4.836986
1002_f_at	3.772763	3.979145	3.529124	3.656052	4.121548	4.146230	3.956283
1003_s_at	6.041962	6.013443	5.156087	5.528908	6.054917	6.028770	5.990144
1004_at	5.751041	6.138876	5.038015	5.396687	5.984467	6.282975	5.860502
1005_at	8.324171	6.379843	8.841847	7.536542	8.685069	9.657075	8.481350
	16007	17003	18001	19002	19008	19014	19017
1000_at	7.675929	7.662426	7.584008	7.840099	7.164922	7.843162	7.695714
1001_at	4.959669	5.743215	4.674920	5.208166	4.554529	5.718569	4.498515
1002_f_at	3.819324	4.142183	3.556962	3.942920	3.541456	4.000303	3.667304
1003_s_at	5.763215	6.307570	5.825432	6.181242	5.869161	6.131560	5.787218
1004_at	5.781565	5.857727	5.786847	5.935438	5.628791	5.613913	5.226277
1005_at	9.314393	10.336505	9.238966	8.941103	8.005422	9.629751	10.907283
	20005	24006	26009	28008	28009	31015	37001
1000_at	7.520867	7.836577	7.470524	7.520806	7.646947	7.727560	7.849455

```

1001_at 5.135697 5.129836 5.213340 4.690815 4.902946 4.866731 4.959450
1002_f_at 4.165152 4.002384 4.126231 3.655414 3.894576 3.970375 4.047428
1003_s_at 6.019615 6.110763 6.271189 5.513344 5.814061 5.779816 6.193790
1004_at 5.850529 5.847115 6.089797 5.211162 5.828099 5.738370 5.824930
1005_at 8.034295 7.967475 9.382400 7.727615 8.067144 9.211820 10.408645
      43006 43015 44001 49004 56007 64005 65003
1000_at 7.960842 8.188617 7.399999 7.813474 7.816922 7.913249 7.800199
1001_at 4.537677 5.154500 5.071885 4.874525 4.788699 5.403640 5.443827
1002_f_at 3.694877 3.949546 3.837517 3.767672 4.248075 4.119252 3.957468
1003_s_at 5.973162 6.033704 5.560886 5.930329 5.878533 6.193376 6.006443
1004_at 5.841095 5.870163 5.686555 5.793403 5.913503 6.053053 5.549087
1005_at 7.524865 8.381740 8.055760 7.234242 8.083834 10.163355 8.184442
      83001 LAL4
1000_at 8.030047 7.702217
1001_at 5.178633 5.029670
1002_f_at 4.053458 4.002306
1003_s_at 6.050129 6.297549
1004_at 5.932412 5.827131
1005_at 8.798521 7.687370

```

Los datos fenotípicos los tendremos con `Biobase::pData`.

```
pData(ALL)
```

¿Cuál fue chip que se utilizó para obtener estos datos?

```
annotation(ALL)
```

```
[1] "hgu95av2"
```

Los identificadores de las filas los tenemos con

```
featureNames(ALL)
```