

(a)

Caryoscope settings

Chromosome axis

☒

Feature tooltip expression

GenBank \${GBACC} - \${GENE_NAME}

Feature URL expression

http://genome-www5.stanford.edu/cgi-bin/source

Caryoscope

X: 100%

Y: 100%

3395-2004-04-04-2...

1

2

3

4

5

6

7

8

9

10

11

12

13

14

15

H47714 - major histocompatibility complex, class II, D

SOURCE

GeneReport

H. sapiens

HLA-DRA

SOURCE Search : (Source::GenePage for Gene H47714)

+

http://genome-www5.stanford.edu/cgi-bin/source/s

Google

Switch to CloneReport

mouse pointer

click navigates to URL

(b)

	Expression	Result
Feature tooltip	GenBank \${GBACC} - \${GENE_NAME}	GenBank H47714 - major histocompatibility co...
Feature URL	http://genome-...&criteria= \${GBACC} &choice=Gene	http://genome-...&criteria= H47714 &choice=Gene

SEQ	FEATURE	START	END	GBACC	CH2I_MEAN	LOG_RAT2N_MEAN	CH2D_MEAN	GENE_NAME
5	clone	35588770	35589227	H11369	731	0.425	81	***START domain containing
6	clone	35588770	35589227	W86653	649	0.116	443	
6	clone	30018888	30019496	AA644657	680	0.15	458	
6	clone	32482382	32483756	H47714	1293	0.243	1120	"major histocompatibility c
8	clone	123854254	123854613	R10310	1596	2.799	1345	"major histocompatibility c
11	clone	109639378	109639788	AA479781	857	0.492	594	"CDNA FLJ27381 fis, clone
12	clone	94390321	94400349	AA488716	1097	-0.267	873	radixin
16	clone	61598469	61626818	AA488716	679	-0.236	464	methionyl aminopeptidase 2
17	clone	38436158	38437150	R56301	778	-0.183	602	"cadherin 8, type 2"
17	clone	38436158	38437150	N45441	4401	2.662	4147	gasdermin-like
20	clone	46590096	46590406	R11529	2049	2.051	1786	