

Chromosome	Number of Genes
1	2451
2	1167
3	1791
4	1054
5	984
6	1295
7	708
8	749
9	1025
10	490
11	588
12	2562
13	2098
14	1900
15	2221
16	2328
17	3499
18	1690
19	1961
20	2030
21	1567

Number of genes

Category	Number of genes
2	2158
3	43
4	11
5	1
6	19
7	3
8	14
9	4
10	2
11	218
12	55
13	1
14	3965
15	2202
16	564
17	69
18	6223
19	568
20	8725
21	78
22	22
23	354
24	51
25	1
26	251
27	24
28	1508
29	1383
30	2
31	2559
32	129
33	6
34	161
35	6
36	4
37	70
38	10
39	144
40	34
41	22
42	239

Chromosome	Percentage
chr1	71.5%
chr10	62.7%
chr11	59.5%
chr12	68.5%
chr13	64.7%
chr14	71.9%
chr15	55.8%
chr16	62.9%
chr17	59.6%
chr18	54.6%
chr19	54.4%
chr2	67.5%
chr3	72.7%
chr4	66.2%
chr5	68.4%
chr6	74.4%
chr7	74.2%
chr8	66.6%
chr9	68.6%
chrX	77.7%
chrY	99.8%

fraction of total genes

Genomic Feature	Fraction of Total Genes (%)
3' _overlapping_ncRNA	100.0%
antisense_intronic	9.2%
IG_C_gene	0.5%
IG_D_gene	84.6%
IG_J_gene	100.0%
IG_L_gene	100.0%
IG_T_gene	100.0%
IG_V_gene	100.0%
IG_V_pseudogene	100.0%
intron	100.0%
macro_intron	3.2%
miRNA	100.0%
misc_RNA	100.0%
polymorphic_pseudogene	9.6%
processed_transcript	85.4%
protein_coding	5.4%
pseudogene	78.8%
ribozyme	9.7%
rRNA	100.0%
scaRNA	100.0%
scRNA	100.0%
sense_intronic	100.0%
sense_overlapping_snoRNA	96.0%
snoRNA	3.0%
snRNA	100.0%
sRNA	100.0%
TCC	25.0%
transcribed_processed_pseudogene	5.5%
transcribed_unitary_pseudogene	75.0%
TR_C_gene	84.3%
TR_D_gene	5.0%
TR_J_gene	100.0%
TR_V_pseudogene	100.0%
unitary_pseudogene	100.0%
unprocessed_pseudogene	4.6%