

Supplemental Table S3. Genomic features present on ENCODE tiling microarray

Region	Chr	Size (Mb)	Transcripts	Genes	CpG Islands	Genic (Mb)	Non-genic (Mb)
ChrX Pick	X	1.2	101	57	47	0.83	0.51
Interleukin	5	1	39	24	21	0.44	0.56
<i>CFTR</i>	7	1.9	23	14	9	0.84	1.04
HOXA Cluster	7	0.5	20	16	39	0.05	0.48
<i>FOXP2</i>	7	1	12	3	1	0.37	0.63
Manual A	7	1.1	13	7	6	0.16	0.95
Manual B	7	1.2	8	6	5	0.84	0.33
<i>APO</i> Cluster	11	0.5	9	7	6	0.04	0.46
<i>HBB</i>	11	1	53	45	0	0.16	0.84
<i>IGF2/H19</i>	11	0.6	38	21	28	0.13	0.48
<i>HBA1</i>	16	0.5	30	23	39	0.28	0.22
Chr19 region	19	1	75	42	36	0.48	0.52
Chr21 region	21	1.7	42	25	23	0.79	0.91
Chr22 region	22	1.7	33	19	14	1.21	0.49
Total	--	14	496	309	274	6.6	8.3