



Table with 40 columns of gene identifiers and numerical values. The table lists various genes such as 19H14.DMA, 19H14.DMA, 19H14.DMA, etc., with corresponding values in each column.

Table with 4 columns: ID, Name, Value, and another Name. The table lists various identifiers and their corresponding values, organized in a grid-like structure.

Table with 3 columns: Gene symbol (e.g., 500 NAKM2), Gene coordinates (e.g., 100 001 55182), and Gene function (e.g., 050 UGC2B). The table lists numerous genes and their genomic coordinates and functions.

Table with 4 columns: Gene Symbol, RefSeq ID, RefSeq Description, and RefSeq Length. The table lists numerous genes and their associated identifiers, such as 778 NMT1, 778 NMT2, 778 NMT3, etc., up to 908 PDLIM4.

Table with 10 columns of alphanumeric codes and numbers, representing a large dataset of identifiers and values.

Table with 4 columns: Gene ID, Gene Name, Chromosome, and Position. The table lists numerous genes such as 116R PSMAB, 116R PSMAC, 116R PSMAD, etc., along with their genomic coordinates.

Table with 4 columns: ID, Gene Symbol, Gene Name, and Value. The table lists numerous genes and their associated values, such as 1360 CHW139, 1361 SLC25A3, 1362 TLR2, etc.



Table with 4 columns: Gene Symbol, X Chromosome Position (Mb), Y Chromosome Position (Mb), and Gene Name. The table lists numerous genes and their genomic coordinates, such as 1568 WARS2, 1568 TRKB, 1568 MTRF1, etc., up to 1748 GSDMC2.





Table with 4 columns: ID, X, Y, Z. Contains a dense list of alphanumeric identifiers and numerical values, likely representing a coordinate system or data points for a 3D model.

Table with 4 columns: ID, Name, Value, and Unit. The table lists various identifiers and their corresponding values and units, such as 2381 ADH1141-1547, 2381 ADH1141-1547, 2381 ADH1141-1547, etc.

Table with 10 columns: ID, X1, X2, X3, X4, X5, X6, X7, X8, X9. Contains a dense grid of alphanumeric characters representing a dataset or codebook.

Table with 4 columns: Gene Symbol, Gene Name, RefSeq ID, and coordinates. The table lists numerous genes such as 2725 S2N, 2726 INKDC2, 2727 AD101B1, etc., with their corresponding identifiers and genomic coordinates.

Table with 100 columns of alphanumeric codes and values, representing a large dataset of identifiers and their associated numerical values.



Table with 4 columns: ID, X1, X2, X3. Contains a dense list of alphanumeric identifiers and numerical values, likely representing a dataset or a list of items with associated coordinates or weights.

Table with 10 columns: Gene ID, Gene Name, Chr, Start, End, Strand, RefSeq, Ensembl, UniProt, and other identifiers. The table lists numerous genes such as 3316 RAG2B, 3317 RAG2, 3318 RAG3, etc., with their genomic coordinates and associated identifiers.

Table with multiple columns containing alphanumeric codes and numerical values. The codes are organized in groups, such as 3509, 3510, 3511, etc., with sub-headers like 3509 9R9L2, 3510 9R9L1, etc. Each entry consists of a code followed by a series of numbers and letters, representing a specific identifier and its associated data points.

Table with 4 columns: ID, Gene, Value, Gene, Value, Gene, Value, Gene, Value. Contains a dense list of gene names and associated numerical values.

Table with 4 columns: ID, X, Y, Z. Contains a dense grid of numerical data points for various identifiers.

Table with 10 columns: Gene ID, Gene Name, RefSeq ID, RefSeq Name, RefSeq Description, RefSeq Accession, RefSeq Date, RefSeq Version, RefSeq Type, RefSeq Status. The table lists numerous genes and their associated identifiers.

Table with multiple columns containing alphanumeric codes (e.g., 4268 G6T4, 4268 H8A2) and numerical values. The table is organized in a grid-like structure with approximately 10 columns per row.

Table with 4 columns: ID, X, Y, Z. Contains a dense grid of alphanumeric characters representing a coordinate system or data points.



Table with multiple columns of alphanumeric codes and numerical values, likely representing a list of identifiers or data points.

Table with 4 columns: Gene ID, Gene Name, RefSeq ID, and RefSeq Name. The table lists numerous genes and their corresponding identifiers, such as 4870 HNRN, 4871 SFRP, 4872 WDR5, etc., up to 9085 CYP27B1.

Table with 10 columns containing alphanumeric codes and numerical values. The table lists various identifiers such as S068 UC7N, S068 UC7A, S068 UC7B, etc., followed by their corresponding numerical values across multiple columns.

Table with 10 columns of alphanumeric codes (e.g., S060 P030M, S061 C0001T1, S062 L12, etc.) and corresponding numerical values.

Table with 4 columns: ID, X, Y, Z. Contains a dense grid of alphanumeric characters representing a dataset or coordinate system.

Table with multiple columns containing alphanumeric codes and numerical values. The table is organized into several columns, with the first column being the most prominent and containing the majority of the data points.

Table with 10 columns: Gene ID, Gene Name, Chr, Start, End, Strand, RefSeq, Ensembl, UniProt, RefSeq, Ensembl, UniProt. The table lists numerous genes and their genomic coordinates across various chromosomes.

Table with 10 columns: ID, X1, X2, X3, X4, X5, X6, X7, X8, X9. Contains a dense grid of alphanumeric codes and values.



Table with 4 columns: ID, X, Y, Z. Contains a dense grid of alphanumeric characters representing a coordinate system or data points.

Table with 10 columns: Gene ID, Gene Name, RefSeq ID, RefSeq Name, RefSeq Description, RefSeq Accession, RefSeq Date, RefSeq Status, RefSeq Type, RefSeq Category. The table lists numerous genes and their associated identifiers.

Table with 4 columns: ID, X, Y, Z. Contains a dense grid of alphanumeric characters representing a coordinate system or data points.

Table with 10 columns of alphanumeric codes and values, representing a large dataset of identifiers and their corresponding numerical values.

Table with 4 columns: ID, X1, X2, X3. Contains a dense list of alphanumeric identifiers and numerical values, likely representing a dataset or a list of items.

Table with 10 columns: Gene symbol, RefSeq ID, Start, End, Strand, Feature, Start, End, Strand, Feature, Start, End, Strand, Feature. Lists genomic coordinates and features for various genes including ABCA1, ABCA2, ABCA3, ABCA4, ABCA5, ABCA6, ABCA7, ABCA8, ABCA9, ABCA10, ABCA11, ABCA12, ABCA13, ABCA14, ABCA15, ABCA16, ABCA17, ABCA18, ABCA19, ABCA20, ABCA21, ABCA22, ABCA23, ABCA24, ABCA25, ABCA26, ABCA27, ABCA28, ABCA29, ABCA30, ABCA31, ABCA32, ABCA33, ABCA34, ABCA35, ABCA36, ABCA37, ABCA38, ABCA39, ABCA40, ABCA41, ABCA42, ABCA43, ABCA44, ABCA45, ABCA46, ABCA47, ABCA48, ABCA49, ABCA50, ABCA51, ABCA52, ABCA53, ABCA54, ABCA55, ABCA56, ABCA57, ABCA58, ABCA59, ABCA60, ABCA61, ABCA62, ABCA63, ABCA64, ABCA65, ABCA66, ABCA67, ABCA68, ABCA69, ABCA70, ABCA71, ABCA72, ABCA73, ABCA74, ABCA75, ABCA76, ABCA77, ABCA78, ABCA79, ABCA80, ABCA81, ABCA82, ABCA83, ABCA84, ABCA85, ABCA86, ABCA87, ABCA88, ABCA89, ABCA90, ABCA91, ABCA92, ABCA93, ABCA94, ABCA95, ABCA96, ABCA97, ABCA98, ABCA99, ABCA100.

Table with 4 columns: Gene ID, Gene Name, RefSeq ID, and RefSeq Name. The table lists numerous genes such as 7405 C12orf66, 7406 C12orf67, 7407 C12orf68, etc., up to 7598 LOC101928. Each entry includes a numerical ID, a gene symbol, and its corresponding RefSeq identifier and name.

Table with 4 columns: Gene ID, Gene Name, RefSeq ID, and RefSeq Name. This table continues the list of genes from the previous table, starting with 7599 LOC101929 and ending with 7600 LOC101930. It follows the same format of Gene ID, Gene Name, RefSeq ID, and RefSeq Name.

Table with 5 columns: Gene ID, Gene Name, RefSeq ID, RefSeq Description, and Gene Location. It lists numerous genes such as 7601 FH4, 7601 FH5, 7601 GBI, etc., with their corresponding identifiers and genomic coordinates.



Table with 4 columns: Gene ID, Gene Name, RefSeq ID, and RefSeq Name. The table lists numerous genes such as 7786 RBM10, 7786 RBM11, 7786 RBM12, etc., with their corresponding identifiers and names.



Table with 10 columns: Gene ID, Gene Name, RefSeq ID, RefSeq Description, Gene ID, Gene Name, RefSeq ID, RefSeq Description, Gene ID, Gene Name, RefSeq ID, RefSeq Description. The table lists numerous genes and their associated identifiers.

Table with 4 columns: Gene ID, Gene Name, RefSeq ID, and coordinates. Lists various genes such as 8301 IER1, 8302 IER2, 8303 IER3, etc., with their corresponding RefSeq IDs and genomic coordinates.

Table with 4 columns: ID, Name, Value, and another Value. The table contains a dense list of alphanumeric identifiers and their corresponding numerical values, organized in a grid-like structure.



Table with 10 columns: ID, X, Y, Z, W, V, U, T, S, R. Contains a dense grid of alphanumeric characters representing genomic coordinates and identifiers.

A large table containing a list of genetic markers, each with a unique ID, a name, and associated numerical values (e.g., -0.44, 0.00, 0.01, etc.). The table is organized in columns and contains thousands of rows of data.



Table with 4 columns: Gene Symbol, chr, start, end, Gene Symbol, chr, start, end, Gene Symbol, chr, start, end, Gene Symbol, chr, start, end. It lists genomic coordinates for various genes across chromosomes 1-22.

Table with 4 columns: ID, Name, Value, and another Value. The table lists numerous identifiers and their corresponding values, organized in a grid-like structure.

Table with 10 columns: Gene ID, Gene Name, Value 1, Value 2, Value 3, Value 4, Value 5, Value 6, Value 7, Value 8. The table lists numerous genes such as 9748 SLC18B3, 9749 SLC16A9, 9750 SLC16A8, etc., with associated numerical values.

Table with 4 columns: ID, X1, X2, X3. Contains a dense list of alphanumeric identifiers and numerical values.

Table with 3 columns: Gene ID, Gene Name, and Value. The table lists numerous genes such as 101813, 101814, 101815, etc., with their corresponding values.

10300 WJF3	-021	010	STK0A	-448	000	ADG5L1	-143	000	EC70A	-1843	000	1980	000	SC129A	339	000	1980C2A	020	010	GGNF-AS1	-1248	000	1980D0A	-118	000	PGC2	-243	022
10301 ZCZC3	-074	014	WV11	-129	004	WV11	-143	000	FN031	-1843	000	1980	000	FMAB02A	332	000	1980F0A	-037	010	AD0029E2	-1319	000	1980G0A	-039	010	DC07	-049	022
10302 X0D1	-178	010	MAKRA	-094	000	TRF0R	-407	000	ZB703	-1843	000	1980	000	SID047	361	000	1980P01	-102	010	ACT10R	-1147	000	1980R0A	-070	000	GNL3L	-049	022
10303 M0P5S	-047	007	EM029	-044	000	EM029	-044	000	EM029	-1843	000	1980	000	AD0029E2	244	000	1980Q01	-037	010	ACT10R	-1147	000	1980S0A	-070	000	GNL3L	-049	022
10304 M1K31	-185	001	LMBR02	-072	000	PTFCR	-449	000	BB025	-1843	000	1980	000	MEM028	157	000	1980T01	-201	010	STAF0A9	-1107	000	1980U0A	-038	000	PNF08L	-039	022
10305 W0M02	-170	003	W0M02	-170	003	W0M02	-170	003	W0M02	-1843	000	1980	000	W0M02	157	000	1980V01	-201	010	STAF0A9	-1107	000	1980W0A	-038	000	PNF08L	-039	022
10306 PALM0	-427	001	MA008	-030	000	AP00709	-055	000	LR054	-1843	000	1980	000	ZNF10	-001	010	1980X01	-001	010	MA008	-030	000	1980Y0A	-038	000	FU02	-334	022
10307 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1980Z01	-201	010	STAF0A9	-1107	000	198100A	-038	000	PNF08L	-039	022
10308 M0P5S	-427	001	MA008	-030	000	AP00709	-055	000	LR054	-1843	000	1980	000	ZNF10	-001	010	1981101	-001	010	MA008	-030	000	198120A	-038	000	FU02	-334	022
10309 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1981301	-001	010	MA008	-030	000	198140A	-038	000	FU02	-334	022
10310 X0D1	-178	010	MAKRA	-094	000	TRF0R	-407	000	ZB703	-1843	000	1980	000	SID047	361	000	1981501	-001	010	MA008	-030	000	198160A	-038	000	FU02	-334	022
10311 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1981701	-001	010	MA008	-030	000	198180A	-038	000	FU02	-334	022
10312 X0D1	-178	010	MAKRA	-094	000	TRF0R	-407	000	ZB703	-1843	000	1980	000	SID047	361	000	1981901	-001	010	MA008	-030	000	198200A	-038	000	FU02	-334	022
10313 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1982101	-001	010	MA008	-030	000	198220A	-038	000	FU02	-334	022
10314 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1982301	-001	010	MA008	-030	000	198240A	-038	000	FU02	-334	022
10315 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1982501	-001	010	MA008	-030	000	198260A	-038	000	FU02	-334	022
10316 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1982701	-001	010	MA008	-030	000	198280A	-038	000	FU02	-334	022
10317 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1982901	-001	010	MA008	-030	000	198300A	-038	000	FU02	-334	022
10318 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1983101	-001	010	MA008	-030	000	198320A	-038	000	FU02	-334	022
10319 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1983301	-001	010	MA008	-030	000	198340A	-038	000	FU02	-334	022
10320 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1983501	-001	010	MA008	-030	000	198360A	-038	000	FU02	-334	022
10321 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1983701	-001	010	MA008	-030	000	198380A	-038	000	FU02	-334	022
10322 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1983901	-001	010	MA008	-030	000	198400A	-038	000	FU02	-334	022
10323 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1984101	-001	010	MA008	-030	000	198420A	-038	000	FU02	-334	022
10324 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1984301	-001	010	MA008	-030	000	198440A	-038	000	FU02	-334	022
10325 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1984501	-001	010	MA008	-030	000	198460A	-038	000	FU02	-334	022
10326 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1984701	-001	010	MA008	-030	000	198480A	-038	000	FU02	-334	022
10327 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1984901	-001	010	MA008	-030	000	198500A	-038	000	FU02	-334	022
10328 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1985101	-001	010	MA008	-030	000	198520A	-038	000	FU02	-334	022
10329 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1985301	-001	010	MA008	-030	000	198540A	-038	000	FU02	-334	022
10330 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1985501	-001	010	MA008	-030	000	198560A	-038	000	FU02	-334	022
10331 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1985701	-001	010	MA008	-030	000	198580A	-038	000	FU02	-334	022
10332 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1985901	-001	010	MA008	-030	000	198600A	-038	000	FU02	-334	022
10333 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1986101	-001	010	MA008	-030	000	198620A	-038	000	FU02	-334	022
10334 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1986301	-001	010	MA008	-030	000	198640A	-038	000	FU02	-334	022
10335 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1986501	-001	010	MA008	-030	000	198660A	-038	000	FU02	-334	022
10336 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1986701	-001	010	MA008	-030	000	198680A	-038	000	FU02	-334	022
10337 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1986901	-001	010	MA008	-030	000	198700A	-038	000	FU02	-334	022
10338 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1987101	-001	010	MA008	-030	000	198720A	-038	000	FU02	-334	022
10339 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1987301	-001	010	MA008	-030	000	198740A	-038	000	FU02	-334	022
10340 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1987501	-001	010	MA008	-030	000	198760A	-038	000	FU02	-334	022
10341 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1987701	-001	010	MA008	-030	000	198780A	-038	000	FU02	-334	022
10342 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1987901	-001	010	MA008	-030	000	198800A	-038	000	FU02	-334	022
10343 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1988101	-001	010	MA008	-030	000	198820A	-038	000	FU02	-334	022
10344 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1988301	-001	010	MA008	-030	000	198840A	-038	000	FU02	-334	022
10345 W0K11	-143	003	W0K11	-143	003	W0K11	-143	003	W0K11	-1843	000	1980	000	W0K11	157	000	1988501	-001	010	MA008	-030	000	198860A	-038	000	FU02	-334	022
10346 W0K11	-143	003	W0K11	-143	003	W0K11																						



Table with 4 columns: Gene symbol, Gene name, Chromosome, and coordinates. The table lists numerous genes such as LOC101928, LOC101929, LOC101930, etc., with their corresponding genomic locations.



Table with 4 columns: Gene Name, RefSeq ID, Start, End, Strand, and Coordinates. Lists numerous genes and their genomic coordinates across various chromosomes.

Table with 4 columns: ID, Gene, Value, and another Value. The table lists numerous genes such as 11101 CBR4, 11101 CBR2, 11101 CBR1, etc., with associated numerical values.

11308 T89A0	-442	0.00	GAMT	-024	0.00	SLC26A11	-035	0.00	CLDN8	-039	0.00	SOX1-AS1	478	0.00	IT7	479	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11308 C89E2	-102	0.00	SLC6A6	-021	0.00	UCAT1	-215	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11308 F8E2	-103	0.00	PRMT2	-012	0.00	LUMI1A	-192	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11308 TRAU	-447	0.00	PRMT2	-012	0.00	LUMI1A	-192	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11311 T18A	-024	0.00	NDCD1	-095	0.00	MTOR	-010	0.00	CD46	-014	0.00	ITEL-AS1	199	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11312 T89D	-054	0.00	PRMT2	-012	0.00	LUMI1A	-192	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11313 O3A	-022	0.00	ROR	-047	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11314 F89K	-093	0.00	PRMT2	-012	0.00	LUMI1A	-192	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11315 A82R1	-482	0.00	HGCG18	-018	0.00	DNAH9	034	0.00	GRR	-251	0.00	LEF2A2	358	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11316 M121A	-019	0.00	POU3	-090	0.00	ADAM14	-136	0.00	DCAMK1-AS1	-030	0.00	SLC35A1	478	0.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11317 A3T5D	-010	0.00	HNF4B1	-010	0.00	ZNF484	-211	0.00	HSTH2PH8	-077	0.00	SLC35A1	478	0.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11318 N10D12	-040	0.00	DNAH9	-034	0.00	ITFC2	-348	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11319 S99A12	-140	0.00	AT	-377	0.00	C17orf132	-045	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11320 T18A	-122	0.00	ROR	-047	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11321 F8E4	-045	0.00	PRMT2	-012	0.00	LUMI1A	-192	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11322 F9A4	-102	0.00	LRG1	-442	0.00	PRMT2	-012	0.00	HNRH3-AS1	-196	0.00	ITFC2	-348	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00
11323 O3A	-022	0.00	ROR	-047	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11324 M3A	-102	0.00	MMP1	-241	0.00	CD46	-014	0.00	FAM189A	-186	0.00	ZNF484	-211	0.00	HSTH2PH8	-077	0.00	SLC35A1	478	0.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00
11325 C2D4A	-188	0.00	HNF1	-150	0.00	HNF1A	-354	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11326 C2D4C14	-015	0.00	DUXO3	-131	0.00	GATB	-111	0.00	BDNF	-199	0.00	ZNF484	-211	0.00	HSTH2PH8	-077	0.00	SLC35A1	478	0.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00
11327 M8C0	-153	0.00	HNF1	-150	0.00	HNF1A	-354	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11328 TAF1	-050	0.00	MAMLI1	-050	0.00	FUT11	-111	0.00	SLC43A	-004	0.00	RHOA2	381	0.00	ITFC2	-348	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00
11329 A48C3C1	-077	0.00	SMO6B	-046	0.00	MMX2	-050	0.00	ZNF71	027	0.00	ZNF484	-211	0.00	HSTH2PH8	-077	0.00	SLC35A1	478	0.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00
11330 D2C1	-138	0.00	HNF4	008	0.00	IPAK	-063	0.00	TBRX8	013	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00
11331 C13C	-108	0.00	JAMMP2	-127	0.00	ANKRD6	-139	0.00	ANKRD6	-139	0.00	CRO73	-033	0.00	RFC6	-147	0.00	PTPFR14	-484	0.00	CFHR4	-048	0.00	DMG1	-042	0.00	SDS24C1	-219	0.00
11332 F89D	-112	0.00	THRAP3	-033	0.00	PALCA	-151	0.00	SPAST	035	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00
11333 M8D8	-168	0.00	ZK	-014	0.00	ABHD13	030	0.00	ABCA3	-035	0.00	FANC8	004	0.00	CNTF	-157	0.00	MACROD2	-281	0.00	KIF3B3	-182	0.00	AP5L2	-030	0.00	FOXA1	-028	0.00
11334 F89L4	-058	0.00	DRE1	-033	0.00	NARS2	-083	0.00	BE89	04	0.00	UTS1B2	843	0.00	MORC4	-168	0.00	RBM4	-012	0.00	NDUF2-AS2	-158	0.00	HTBA2	042	0.00	PGC4	-024	0.00
11335 F89L5	-157	0.00	DRG2	-237	0.00	PPP2R3	002	0.00	ACOR4A	-221	0.00	HIF1F3	287	0.00	SPY5	-157	0.00	FRS2	-017	0.00	NDUF2-AS2	-158	0.00	HTBA2	042	0.00	PGC4	-024	0.00
11336 F89L6	-049	0.00	HAZ1	-016	0.00	ABCF3	000	0.00	AMC58	027	0.00	CERAM	-028	0.00	TRX2	-152	0.00	CEACCP2	-078	0.00	GRP3C2	-180	0.00	PHK1	-011	0.00	INAM2	-151	0.00
11337 F89L7	-019	0.00	HNF1O1	-013	0.00	PCNA	010	0.00	TWAP	026	0.00	ZNF484	-211	0.00	HSTH2PH8	-077	0.00	SLC35A1	478	0.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00
11338 F89L8	-049	0.00	HAZ1	-016	0.00	ABCF3	000	0.00	AMC58	027	0.00	CERAM	-028	0.00	TRX2	-152	0.00	CEACCP2	-078	0.00	GRP3C2	-180	0.00	PHK1	-011	0.00	INAM2	-151	0.00
11339 F89L9	-049	0.00	HAZ1	-016	0.00	ABCF3	000	0.00	AMC58	027	0.00	CERAM	-028	0.00	TRX2	-152	0.00	CEACCP2	-078	0.00	GRP3C2	-180	0.00	PHK1	-011	0.00	INAM2	-151	0.00
11340 D2C2B	-058	0.00	RARL	-053	0.00	HSPB1	-027	0.00	RLR2	-002	0.00	ZNF484	-211	0.00	HSTH2PH8	-077	0.00	SLC35A1	478	0.00	IT7-AS2	478	0.00	CCAT7	-141	0.00	NR2F36	-174	0.00
11341 F89L10	-140	0.00	FAM189A	-186	0.00	ITFC2	-348	0.00	NOTD	-029	1.00	IT7-AS2	478	0.00	ANGPT4	-132	0.00	SRM8A	-177	0.00	NR2F36	-174	0.00	MMMP9	-190	0.00	CHMR2	-031	0.00
11342 F89L11	-039	0.00	THAP3	-033	0.00	PALCA	-151	0.00	SPAST	035	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00	UNC53B	422	0.00
11343 F89L12	-148	0.00	HNF4	008	0.00	IPAK	-063	0.00	TBRX8	013	0.00	CRO73	-033	0.00	RFC6	-147	0.00	PTPFR14	-484	0.00	CFHR4	-048	0.00	DMG1	-042	0.00	SDS24C1	-219	0.00
11344 F89L13	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11345 F89L14	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11346 F89L15	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11347 F89L16	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11348 F89L17	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11349 F89L18	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11350 F89L19	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11351 F89L20	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11352 F89L21	-029	0.00	DMO	-064	0.00	ZNF703	047	0.00	TMEM127	-044	0.00	HELI1	068	0.00	NR6	-016	0.00	MCE1	-014	0.00	CASC20	-190	0.00	CSC21	-190	0.00	GCAT	-159	0.00
11353 F89L22	-029	0.00	DMO	-0																									

Table with multiple columns containing alphanumeric codes and values. The table lists various identifiers such as 1508 HRCB, 1508 HRCB, 1508 HRCB, etc., followed by numerical values and corresponding codes like 0001, 0002, 0003, etc.

Table with 10 columns: Gene ID, Gene Name, Chr, Start, End, Strand, RefSeq, RefSeq ID, RefSeq Description, RefSeq Accession. Contains a comprehensive list of genes and their genomic coordinates across various chromosomes.

Table with 4 columns: Gene ID, Gene Name, and two columns of numerical values. The table lists numerous genes such as 1880 TMEM254, 1881 CDC2, 1882 TRAF6, etc., with associated values.

Supplemental material table with columns for gene names and numerical values. The table is organized into columns of approximately 10-15 genes each, with values ranging from -0.99 to 0.99. The genes listed include various human genes such as C11orf105, GNAK2, ABCB9, CASA, and many others.

Table with 10 columns: Gene ID, Gene Name, RefSeq ID, RefSeq Name, RefSeq Description, RefSeq Accession, RefSeq Date, RefSeq Status, RefSeq Type, RefSeq Category. The table lists numerous genes and their associated identifiers.



12475 AVEH	-105	00	ZC9H	-032	00	MFA3	-094	00	WDC1	-338	00	YFC2B	292	00	MIK2	-098	00	MMP17	-089	00	RAVW	-081	04	FAD2	-153	00	FOA1M	-239	00	02
12476 KZM4	-127	00	VCM3	-187	00	CFY4	-074	00	WDC2	209	00	ANW2B	321	00	MIK3	-037	00	WDC3	-122	00	WDC4	-138	04	WDC5	-029	00	SC3M4	-334	00	02
12477 A2A	-105	00	ATP3A	-126	00	FNZ3	-415	00	TME213	330	00	URC2	392	00	CXK1	-023	00	CAP15	-184	00	AD2M17	-189	04	SHG8	-442	00	PPN2C	-043	00	02
12478 WZKAP	-055	00	AM3	-034	00	Q6	-016	00	Q6	014	00	URC3	388	00	URC4	-016	00	URC5	-184	00	URC6	-184	00	URC7	-184	00	URC8	-184	00	02
12479 TME9	-168	00	GC2	-022	00	SHZ9	-079	00	BKA	-030	00	TAN2C	-032	00	URC9	-023	00	CIS2	-037	00	ZPW74	-135	04	SHG18	-515	00	AWCA	-171	00	02
12480 WZKAP	-055	00	AM3	-034	00	Q6	-016	00	Q6	014	00	URC3	388	00	URC4	-016	00	URC5	-184	00	URC6	-184	00	URC7	-184	00	URC8	-184	00	02
12481 MFC2	-128	00	UP	-414	00	HAK3A	-104	00	TAF3A	-278	00	URF4	459	00	URF5	-035	00	URF6	-035	00	URF7	-035	00	URF8	-035	00	URF9	-035	00	02
12482 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12483 M72	-105	00	NT56	-005	00	CH4L	-078	00	CH4D	-030	00	URF4	459	00	URF5	-035	00	URF6	-035	00	URF7	-035	00	URF8	-035	00	URF9	-035	00	02
12484 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12485 SC3M4	-083	00	APM1	-015	00	TMC2	-403	00	DAR2	-030	00	URF4	459	00	URF5	-035	00	URF6	-035	00	URF7	-035	00	URF8	-035	00	URF9	-035	00	02
12486 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12487 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12488 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12489 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12490 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12491 MPM2	-259	00	PAR70	-039	00	C1K2	-068	00	URF19	-035	00	URF20	-035	00	URF21	-035	00	URF22	-035	00	URF23	-035	00	URF24	-035	00	URF25	-035	00	02
12492 MPM2	-259	00	PAR70	-039	00	C1K2	-068	00	URF19	-035	00	URF20	-035	00	URF21	-035	00	URF22	-035	00	URF23	-035	00	URF24	-035	00	URF25	-035	00	02
12493 MPM2	-259	00	PAR70	-039	00	C1K2	-068	00	URF19	-035	00	URF20	-035	00	URF21	-035	00	URF22	-035	00	URF23	-035	00	URF24	-035	00	URF25	-035	00	02
12494 MPM2	-259	00	PAR70	-039	00	C1K2	-068	00	URF19	-035	00	URF20	-035	00	URF21	-035	00	URF22	-035	00	URF23	-035	00	URF24	-035	00	URF25	-035	00	02
12495 C2K2B	-022	00	C1K2	-032	00	YOC	-017	00	COCA	-057	00	URF19	-035	00	URF20	-035	00	URF21	-035	00	URF22	-035	00	URF23	-035	00	URF24	-035	00	02
12496 MFC2	-128	00	UP	-414	00	HAK3A	-104	00	TAF3A	-278	00	URF4	459	00	URF5	-035	00	URF6	-035	00	URF7	-035	00	URF8	-035	00	URF9	-035	00	02
12497 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12498 M72	-105	00	NT56	-005	00	CH4L	-078	00	CH4D	-030	00	URF4	459	00	URF5	-035	00	URF6	-035	00	URF7	-035	00	URF8	-035	00	URF9	-035	00	02
12499 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12500 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12501 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12502 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12503 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12504 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12505 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12506 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12507 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12508 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12509 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12510 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12511 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12512 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12513 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12514 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12515 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12516 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12517 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12518 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12519 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12520 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12521 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12522 S2R18	-112	00	URF10	-035	00	URF11	-035	00	URF12	-035	00	URF13	-035	00	URF14	-035	00	URF15	-035	00	URF16	-035	00	URF17	-035	00	URF18	-035	00	02
12523 S2R18	-112	00	URF10	-035																										



18865 TSK1	-129	0.00	STW2	-0.34	0.00	EXV2	-0.66	0.00	ZICW4A	-0.21	0.00	CHN1	2.45	0.00	CELC4C	-4.96	0.00	SICL3A1	-0.43	0.00	GTRHC6	-0.64	0.00	SCS1	-0.19	0.00	DDDKX	0.11	0.00
18866 CDT2	-0.39	0.00	DPW2	0.11	0.00	CSB1E1	2.18	0.00	CHN2	0.29	0.00	CHN1	-0.29	0.00	CELC4C	-4.96	0.00	AMK2	-2.43	0.00	PCCD3C	-1.93	0.00	FCR2B	-1.35	0.00	NAPY1E	-0.04	0.00
18867 TFM	-0.00	0.00	URCS3	-0.00	0.00	KPFC	-0.01	0.00	ZYK	0.07	0.00	GL1	4.93	0.00	NW02	-1.46	0.00	BRM12	-0.46	0.00	COZ	-2.47	0.00	WNV7B	-5.51	0.00	NAPY1E	-0.04	0.00
18868 AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00	AMM1	-0.02	0.00
18869 GMP	-1.15	0.00	EXT1	-1.32	0.00	EXT1	-1.07	0.00	ASC2	0.07	0.00	CHN1	3.41	0.00	CHN1	-0.47	0.00	NK42H	-0.58	0.00	STABDAS	-1.47	0.00	MP	-4.80	0.00	4-SEP	0.30	0.00
18870 MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00	MTA1	-0.22	0.00
18871 C0C71	-0.82	0.00	MU1	-1.22	0.00	DK4P	-0.05	0.00	FAM15A	0.07	0.00	CHN2	1.89	0.00	CHN2	-1.90	0.00	CASK	-1.96	0.00	AR3	-1.17	0.00	SEPH1	-0.40	0.00	PPM1A	0.13	0.00
18872 TFC1	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00	KANP9	-0.65	0.00
18873 M03	-0.00	0.00	KCL2	-0.00	0.00	MTR	-0.54	0.00	OSPBL2	0.24	0.00	LY1	1.17	0.00	ORP2	-0.48	0.00	ORP2	-0.48	0.00	ORP2	-0.48	0.00	ORP2	-0.48	0.00	ORP2	-0.48	0.00
18874 M03	-0.17	0.00	SMK91	-1.83	0.00	SCAMP1	0.21	0.00	UNC0403	-0.22	0.00	UNC0403	-0.22	0.00	UNC0403	-0.22	0.00	UNC0403	-0.22	0.00	UNC0403	-0.22	0.00	UNC0403	-0.22	0.00	UNC0403	-0.22	0.00
18875 F0A2	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00
18876 F0A2	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00
18877 F0A2	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00
18878 F0A2	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00
18879 F0A2	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00	BAGC1	-1.17	0.00
18880 C0C71	-0.82	0.00	MU1	-1.22	0.00	DK4P	-0.05	0.00	FAM15A	0.07	0.00	CHN2	1.89	0.00	CHN2	-1.90	0.00	CASK	-1.96	0.00	AR3	-1.17	0.00	SEPH1	-0.40	0.00	PPM1A	0.13	0.00
18881 S1A	-4.82	0.00	AKC2C	-1.92	0.00	IKB3	-1.21	0.00	TNKS27	0.04	0.00	FAH68B	0.87	0.00	FAH68B	-0.42	0.00	FAH68B	-0.42	0.00	FAH68B	-0.42	0.00	FAH68B	-0.42	0.00	FAH68B	-0.42	0.00
18882 M02	-0.25	0.00	UNC13B	-4.13	0.00	FKBP1B	-0.24	0.00	SIC29A4	-3.11	0.00	STABDAS	-1.47	0.00	MP	-4.80	0.00	MP	-4.80	0.00	MP	-4.80	0.00	MP	-4.80	0.00	MP	-4.80	0.00
18883 S4	-1.65	0.00	UTFC4B	-1.95	0.00	WNA1	-2.18	0.00	SHO2A	-1.05	0.00	GRE1	1.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00
18884 S4	-1.65	0.00	UTFC4B	-1.95	0.00	WNA1	-2.18	0.00	SHO2A	-1.05	0.00	GRE1	1.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00
18885 S4	-1.65	0.00	UTFC4B	-1.95	0.00	WNA1	-2.18	0.00	SHO2A	-1.05	0.00	GRE1	1.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00
18886 S4	-1.65	0.00	UTFC4B	-1.95	0.00	WNA1	-2.18	0.00	SHO2A	-1.05	0.00	GRE1	1.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00	GRE1	-0.25	0.00
18887 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18888 M02	-1.14	0.00	SAPY1	-0.96	0.00	GLCE	-0.83	0.00	CDK3	-0.51	0.00	KCNC1	4.61	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00
18889 M02	-1.14	0.00	SAPY1	-0.96	0.00	GLCE	-0.83	0.00	CDK3	-0.51	0.00	KCNC1	4.61	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00
18890 M02	-1.14	0.00	SAPY1	-0.96	0.00	GLCE	-0.83	0.00	CDK3	-0.51	0.00	KCNC1	4.61	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00
18891 M02	-1.14	0.00	SAPY1	-0.96	0.00	GLCE	-0.83	0.00	CDK3	-0.51	0.00	KCNC1	4.61	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00	KCNC1	-0.23	0.00
18892 SCLG4	-0.74	0.00	MTA1	-0.49	0.00	CDK3	-0.48	0.00	MPD1D	-0.14	0.00	PRB2	3.74	0.00	PRB2	-0.99	0.00	PRB2	-0.99	0.00	PRB2	-0.99	0.00	PRB2	-0.99	0.00	PRB2	-0.99	0.00
18893 M02	-1.01	0.00	CHK3	-0.33	0.00	WNT7B	-4.43	0.00	MSG4B	-0.80	0.00	SHO2	2.75	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00
18894 M02	-1.01	0.00	CHK3	-0.33	0.00	WNT7B	-4.43	0.00	MSG4B	-0.80	0.00	SHO2	2.75	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00
18895 M02	-1.01	0.00	CHK3	-0.33	0.00	WNT7B	-4.43	0.00	MSG4B	-0.80	0.00	SHO2	2.75	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00
18896 M02	-1.01	0.00	CHK3	-0.33	0.00	WNT7B	-4.43	0.00	MSG4B	-0.80	0.00	SHO2	2.75	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00	SHO2	-0.23	0.00
18897 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18898 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18899 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18900 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18901 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18902 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18903 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18904 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18905 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18906 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18907 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18908 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18909 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2	2.02	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00	KCNC2	-0.18	0.00
18910 FAM138	-0.54	0.00	GGP1	0.03	0.00	CDNR	-0.61	0.00	ABC2	-1.87	0.00	KCNC2																	

Table with 3 columns: Gene ID, Gene Name, and Value. The table lists numerous genes such as 13060 TMEM34, 13061 RPL10, 13062 RPL10A, etc., with associated numerical values ranging from -0.18 to 0.24.











Table with 10 columns: Gene ID, Gene Name, Chr, Start, End, Strand, RefSeq, Ensembl, UniProt, and other identifiers. The table lists numerous genes and their genomic coordinates across various chromosomes.









Table with 4 columns: ID, X, Y, Z. Contains a dense grid of numerical data points for various identifiers.



Table with 4 columns: ID, Name, Value, and another Name. The table lists numerous identifiers and their corresponding values, organized in a grid-like structure.



Table with 3 columns: Gene ID, Gene Name, and Correlation Coefficient. The table lists numerous genes such as 15696 TGF1, 15698 TRAF2, 15699 TRAF3, etc., with their respective correlation values.



Table with 100 columns of numerical data representing genomic coordinates and values. The data is organized in a grid-like structure with 100 columns and approximately 1000 rows of data points.



Table with 100 columns of gene names and their corresponding expression values. The table lists various genes such as 16379 SHAR29, 16379 TRAK, 16379 TRAF1, etc., with values ranging from -1.31 to 0.34.

Table with 10 columns of gene names and numerical values. The table lists various genes such as 16070 CROCC1, 16071 EDC1, 16072 RAB18, etc., with corresponding values in each column.

Table with 3 columns: Gene symbol, Gene name, and Value. The table lists numerous genes such as 17K11, 17K12, 17K13, etc., with their corresponding values ranging from -1.16 to 4.20.

Table with 4 columns: Gene ID, Gene Name, Gene Symbol, and Value. The table lists numerous genes such as 19600 TMEM48B, 19601 BMTF, 19602 TMEM160, etc., with associated numerical values.





Table with 10 columns of gene names and numerical values. The table lists various genes such as 1700 DCORL1, 1701 PTPN11, 1702 SLC6A1, etc., with associated numerical values ranging from -130 to 204.

Table with 10 columns of gene symbols and numerical values. The table lists various genes such as 1764 CD47, 1764 CD48, 1764 CD49, etc., with associated numerical values ranging from -0.88 to 0.46.

Table with 4 columns: Gene symbol, Gene name, Chromosome, and Position. The table lists numerous genes such as 17161 C6orf2, 17162 C6orf3, 17163 C6orf4, etc., with their corresponding genomic coordinates.



18130 GPXA	-142	0.00	C37B	-157	0.00	HSPHRI1	-138	0.00	GARG	-295	0.00	EVL	-334	0.00	EY2B	-455	0.00	CTSC	-260	0.00	RP35A	-639	0.00	SERPNA1	-288	0.00	CD3151	-200	0.00
18131 M2FA	-243	0.00	T0N	-137	0.00	AT0D	-189	0.00	BP51	-679	0.00	EY2B	-508	0.00	T0M3	-145	0.00	RP12	-178	0.00	RP12	-165	0.00	PSAP	-150	0.00	MT-CC2	-042	0.00
18132 TH0BP	-322	0.00	S10D0B	-342	0.00	AF1	-341	0.00	CTSC	-219	0.00	PHD5	-089	0.00	APFC	-095	0.00	YWHAB	-108	0.00	RP17A	-042	0.00	HUA-DQ1	-260	0.00	NJRP1	-323	0.00
18133 FL	-125	0.00	HSP91	-217	0.00	HSP91	-166	0.00	KCSE	-292	0.00	FAU	-072	0.00	RP12	-095	0.00	GRF5	-111	0.00	RP12	-056	0.00	SYNGR3	-239	0.00	CC14	-352	0.00
18134 GLU	-351	0.00	ATP8V9B	-187	0.00	KU7A	-346	0.00	TH0BP	-293	0.00	RCC2	-325	0.00	RP12	-054	0.00	EMIA	-208	0.00	RP13	-487	0.00	CD59	-243	0.00	RAB21	-487	0.00
18135 WAPR	-171	0.00	CTSC	-192	0.00	NJRP1	-154	0.00	RP12	-058	0.00	PDOS	-180	0.00	KCSE	-340	0.00	RGCB	-166	0.00	H0PK	-430	0.00	ILN1	-137	0.00	RP130	-396	0.00
18136 HSP91AA1	-119	0.00	CTSC	-110	0.00	S10D43	-317	0.00	CXCA4	-268	0.00	DYFC	-535	0.00	C24B	-456	0.00	SERPNA1	-299	0.00	RP52	-035	0.00	PKM1	-143	0.00	PLAUR	-340	0.00
18137 LGALS3	-157	0.00	S10D411	-121	0.00	CTSC	-159	0.00	EPH2	-159	0.00	RGS1	-444	0.00	CTSC	-245	0.00	ITGB	-182	0.00	TM2C	-482	0.00	CTSC	-143	0.00	LGALS3	-144	0.00
18138 CTSC	-260	0.00	YBC3	-345	0.00	RAB11	-482	0.00	C24B	-383	0.00	EF1A1	-116	0.00	MT-CC1	-045	0.00	CYTP	-187	0.00	RP111	-055	0.00	CTSC	-101	0.00	CER9B	-176	0.00
18139 PDLIM1	-157	0.00	FC02T	-293	0.00	CTSC	-153	0.00	EPH2	-076	0.00	LP1	-447	0.00	CTSC	-061	0.00	BARH52	-182	0.00	RP139	-055	0.00	CTSC	-181	0.00	HSP90B	-123	0.00
18140 HCS7	-239	0.00	GSN	-440	0.00	TMPI1	-191	0.00	RS1	-365	0.00	ITGB	-346	0.00	LP1	-424	0.00	CL1	-079	0.00	RP115	-050	0.00	ANNA2	-147	0.00	AF1	-373	0.00
18141 RPA4	-217	0.00	CTSC	-296	0.00	GLU	-203	0.00	E2F1	-238	0.00	CBAR	-624	0.00	HUA-A	-089	0.00	RP123	-209	0.00	RP52	-058	0.00	HSP90B	-225	0.00	RPA4	-127	0.00
18142 SERPNA1	-149	0.00	SERPNA1	-337	0.00	LGALS3BP	-303	0.00	UBA52	-064	0.00	DP192A2	-223	0.00	TH0BP	-383	0.00	RP138	-051	0.00	RP52	-065	0.00	S10D4B	-096	0.00	POU4F1	-431	0.00
18143 FTH1	-158	0.00	CTSC	-230	0.00	ANNA2	-325	0.00	RP13A	-076	0.00	CTFL1	-309	0.00	CTFL1	-289	0.00	FNAN	-244	0.00	RP18	-062	0.00	YBC3	-375	0.00	CAP5	-391	0.00
18144 ENO3	-164	0.00	PTM5	-295	0.00	CD3151	-230	0.00	RP12	-062	0.00	UBA52	-089	0.00	LRBP1	-207	0.00	RP128	-044	0.00	RP14	-058	0.00	HUA-D8B	-169	0.00	MT-ND3	-059	0.00
18145 LYR6	-184	0.00	DIP	-139	0.00	CTSC	-176	0.00	RP52B	-177	0.00	CDIC4	-400	0.00	CDIC4	-400	0.00	ALS	-187	0.00	RP16	-080	0.00	RCC2	-227	0.00	FECD4	-456	0.00
18146 SGN	-160	0.00	GLU	-258	0.00	CTSC	-219	0.00	YWH0B	-163	0.00	TH0BP	-454	0.00	CTSC	-282	0.00	E2F1	-186	0.00	RP120	-075	0.00	T0N1	-151	0.00	CTSH	-236	0.00
18147 T0N	-187	0.00	TM2P	-114	0.00	SYNGR3	-210	0.00	LP1	-375	0.00	RAC2	-401	0.00	ALDO3AP	-402	0.00	CALM1	-100	0.00	RP12	-056	0.00	GRM1	-152	0.00	RAB13	-404	0.00
18148 RPTM3	-127	0.00	HEAT1	-134	0.00	QSOX1	-129	0.00	QSOX1	-095	0.00	RP52B	-107	0.00	HUA-B	-196	0.00	CC51	-811	0.00	RP127A	-054	0.00	FC02T	-105	0.00	HUA-D9A	-160	0.00
18149 SERP2	-082	0.00	HUA-DQ4	-340	0.00	ALDH2	-364	0.00	RP127A	-067	0.00	RP12	-476	0.00	GSMA3	-400	0.00	COR3D1A	-223	0.00	RP17	-071	0.00	CHCHD10	-124	0.00	S10D4B	-308	0.00
18150 SIZ2	-378	0.00	COX	-315	0.00	RAB13	-423	0.00	RP5A	-101	0.00	RP12	-125	0.00	ARPC3	-143	0.00	MT-CC3	-082	0.00	RP12A	-078	0.00	TMLH1A	-176	0.00	SERPNA1	-296	0.00
18151 PPI1	-122	0.00	ICEG1G	-364	0.00	FCER1G	-276	0.00	MT-CC2	-065	0.00	GRM3	-447	0.00	QAZ1	-098	0.00	CTSC	-209	0.00	RP58	-049	0.00	P1P1	-230	0.00	GSN	-379	0.00
18152 MAT2	-171	0.00	BR1	-196	0.00	SERPNA1	-308	0.00	ALDO3AP	-339	0.00	LP5	-496	0.00	REM	-073	0.00	CTSH	-158	0.00	RP124	-058	0.00	HUA-D4A	-146	0.00	CTSD	-157	0.00
18153 VM	-181	0.00	ASPM1	-259	0.00	CTSC	-295	0.00	RP52B	-089	0.00	COR3D1A	-479	0.00	ITG1	-169	0.00	BR3	-275	0.00	RP225	-048	0.00	PEP1	-181	0.00	APU12	-192	0.00
18154 CL1	-118	0.00	HUA-D9A	-300	0.00	GRM1	-127	0.00	RP129	-080	0.00	LGALS3	-251	0.00	ITGB2	-434	0.00	HUA-D4A	-146	0.00	RP120	-068	0.00	HUA-DQ1	-230	0.00	ALDH2	-439	0.00
18155 CALM1	-138	0.00	SERP2	-079	0.00	S10D4B	-347	0.00	RP12	-086	0.00	CL1	-121	0.00	HUA-D8B1	-251	0.00	HUA-D4A	-125	0.00	RP18	-046	0.00	CTSH	-140	0.00	YBC3	-340	0.00
18156 T0P2	-240	0.00	HUA-DQ8	-378	0.00	TH0BP	-247	0.00	LP1	-370	0.00	CD35	-156	0.00	FN1	-115	0.00	ACTR	-082	0.00	EFSD3	-089	0.00	BR3	-167	0.00	CD41	-086	0.00
18157 ACT1G	-125	0.00	GN	-341	0.00	CD9	-256	0.00	ITGB2	-359	0.00	CXCA4	-577	0.00	COR3D1A	-450	0.00	RCC2	-274	0.00	RP14	-044	0.00	SPN12	-405	0.00	S10D411	-130	0.00
18158 CT3	-296	0.00	LGALS3	-233	0.00	GSN	-405	0.00	MT-CC3	-073	0.00	ACTR	-123	0.00	CD31	-479	0.00	FN1	-102	0.00	RP141	-073	0.00	CD9	-107	0.00	UFAP	-147	0.00
18159 RAC1	-207	0.00	APU12	-109	0.00	CTSH	-280	0.00	S10D4B	-119	0.00	PTPDC	-488	0.00	RP128	-079	0.00	EVL	-154	0.00	RP57	-079	0.00	HUA-D9A	-104	0.00	S10D4B	-094	0.00
18160 CT9	-261	0.00	SAT1	-151	0.00	LGALS3	-184	0.00	RAC2	-065	0.00	HCS7	-502	0.00	RAC2	-470	0.00	SPN12	-186	0.00	RP15	-077	0.00	TM2P	-155	0.00	CD3	-246	0.00
18161 LDHA	-213	0.00	LGALS3	-190	0.00	CDH3	-139	0.00	FTH1	-122	0.00	EMPS	-348	0.00	RP52	-093	0.00	ITGB2	-119	0.00	RP5A	-094	0.00	CPAP	-150	0.00	FC02T	-111	0.00
18162 CSU1	-279	0.00	TH0BP	-369	0.00	SAT1	-184	0.00	LAPTM5	-204	0.00	SH3BP1	-180	0.00	MT-CC2	-070	0.00	HUA-DQ1	-110	0.00	RP51B	-063	0.00	SAT1	-146	0.00	BR1	-247	0.00
18163 LGALS3	-339	0.00	ANNA2	-195	0.00	SPN12	-317	0.00	CD33	-194	0.00	HUA-A	-166	0.00	HUA-D9A1	-295	0.00	LYR6	-187	0.00	RP124	-077	0.00	CAPC	-147	0.00	S10D4D	-120	0.00
18164 ANNA2	-275	0.00	RP12	-226	0.00	FTH1	-108	0.00	RP52	-100	0.00	HUA-E	-191	0.00	PHD5	-234	0.00	CXCA4	-396	0.00	RP121	-067	0.00	CD14	-182	0.00	OST1	-193	0.00
18165 ACT1B	-138	0.00	RAC1	-179	0.00	FC02T	-304	0.00	RP129	-084	0.00	CD37	-502	0.00	PTPDC	-425	0.00	LP1	-430	0.00	RP120	-073	0.00	CD44	-177	0.00	ASPM1	-236	0.00
18166 RZ	-311	0.00	GP2	-224	0.00	RAC1	-148	0.00	COR3D1A	-085	0.00	GARG	-544	0.00	HUA-D9B1	-280	0.00	IL32	-420	0.00	RP15A	-087	0.00	FTH1	-114	0.00	MT-ND4	-059	0.00
18167 TM2B4X	-136	0.00	CTSC	-231	0.00	CTSC	-257	0.00	UFAP	-251	0.00	ITG1	-248	0.00	GARG	-379	0.00	HUA-A	-127	0.00	RP13A	-077	0.00	ANNA2	-147	0.00	GN	-277	0.00
18168 FTH1	-160	0.00	S10D4B	-129	0.00	HSP91	-211	0.00	RP51B	-081	0.00	FN1	-179	0.00	HCS7	-460	0.00	HUA-D9A	-153	0.00	RP13A	-085	0.00	CTSH	-101	0.00	FC02T	-152	0.00
18169 CD83	-225	0.00	HUA-D9B1	-241	0.00	S10D411	-150	0.00	PTPDC	-062	0.00	HUA-B	-181	0.00	CD37	-448	0.00	HUA-C	-141	0.00	RP54X	-094	0.00	PHD5	-104	0.00	PSAP	-171	0.00
18170 S10D43	-104	0.00	PSAP	-241	0.00	ASPM1	-255	0.00	HCS7	-246	0.00	ARHGD3B	-289	0.00	HUA-D9A	-280	0.00	HUA-D9B1	-285	0.00	RP113	-089	0.00	S10D4D	-156	0.00	QUL	-317	0.00
18171 GASH1	-148	0.00	NPC2	-145	0.00	ARPC2	-268	0.00	CD37	-402	0.00	CWA	-243	0.00	CWA	-402	0.00	S10D4D	-189	0.00									