

**Supplementary Table S3: Fine-mapping results from Susie**

| cyto_locus | functional.consequence     | gene.alias    | chr | pos       |
|------------|----------------------------|---------------|-----|-----------|
| bNGF_1     | intron                     | CASC15        | 6   | 22016146  |
| bNGF_1     | intron                     | CASC15        | 6   | 22034854  |
| bNGF_1     | intron                     | CASC15        | 6   | 22060799  |
| bNGF_1     | intron                     | CASC15        | 6   | 22062824  |
| bNGF_1     | intron                     | CASC15        | 6   | 22080016  |
| bNGF_1     | intron                     | CASC15        | 6   | 22085115  |
| bNGF_1     | intron                     | CASC15        | 6   | 22089895  |
| bNGF_1     | intron                     | CASC15        | 6   | 22112043  |
| bNGF_1     | intron                     | CASC15        | 6   | 22113390  |
| bNGF_1     | intron                     | CASC15        | 6   | 22114485  |
| bNGF_1     | intron                     | CASC15        | 6   | 22115797  |
| bNGF_1     | intron                     | CASC15        | 6   | 22122568  |
| bNGF_1     | intron                     | CASC15        | 6   | 22125157  |
| bNGF_1     | intron                     | CASC15        | 6   | 22128962  |
| bNGF_1     | intron                     | CASC15        | 6   | 22130604  |
| bNGF_1     | intron                     | CASC15        | 6   | 22134989  |
| bNGF_1     | non_coding_transcript_exon | NBAT1         | 6   | 22136160  |
| bNGF_1     | non_coding_transcript_exon | NBAT1         | 6   | 22136879  |
| bNGF_1     | intron                     | NBAT1         | 6   | 22140114  |
| bNGF_1     | intron                     | NBAT1         | 6   | 22144059  |
| bNGF_1     | intron                     | CASC15        | 6   | 22208557  |
| bNGF_1     | intron                     | CASC15        | 6   | 22208886  |
| bNGF_1     | intron                     | CASC15        | 6   | 22208958  |
| bNGF_2     | intergenic                 | IGHV3-48      | 14  | 107001019 |
| bNGF_2     | intergenic                 | IGHV3-48      | 14  | 107003217 |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 40969764  |
| EOTAXIN_1  | intergenic                 | RP11-520A21.1 | 3   | 40991176  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41027929  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41039907  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41051489  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41060043  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41075897  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41076810  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41076957  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41081695  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41084811  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41089174  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41100858  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41108552  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41115344  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41118103  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41118898  |
| EOTAXIN_1  | intergenic                 | RP11-944L7.4  | 3   | 41120419  |

|            |            |               |    |          |
|------------|------------|---------------|----|----------|
| EOTAXIN_1  | intergenic | RP11-944L7.4  | 3  | 41137512 |
| EOTAXIN_1  | intergenic | RP11-944L7.4  | 3  | 41140417 |
| EOTAXIN_1  | intergenic | RP11-944L7.4  | 3  | 41146499 |
| EOTAXIN_13 | NA         | NA            | 8  | 59733378 |
| EOTAXIN_13 | NA         | NA            | 8  | 59740982 |
| EOTAXIN_14 | intergenic | LINC01228     | 16 | 79937827 |
| EOTAXIN_14 | intergenic | LINC01228     | 16 | 79951279 |
| EOTAXIN_14 | NA         | NA            | 16 | 79965872 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 79974116 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 79986673 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 79993507 |
| EOTAXIN_14 | NA         | NA            | 16 | 80007176 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80015010 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80019032 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80020499 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80023074 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80034203 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80035498 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80036384 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80042989 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80043989 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80055513 |
| EOTAXIN_14 | NA         | NA            | 16 | 80062214 |
| EOTAXIN_14 | intergenic | RP11-148M9.1  | 16 | 80065251 |
| EOTAXIN_14 | upstream   | RP11-148M9.1  | 16 | 80071830 |
| EOTAXIN_14 | NA         | NA            | 16 | 80089053 |
| EOTAXIN_14 | intergenic | RP11-525K10.1 | 16 | 80093959 |
| EOTAXIN_14 | intron     | RP11-525K10.1 | 16 | 80100004 |
| EOTAXIN_14 | intron     | RP11-525K10.1 | 16 | 80110182 |
| EOTAXIN_14 | intron     | RP11-525K10.1 | 16 | 80112076 |
| EOTAXIN_15 | intron     | RP11-136C24.3 | 3  | 42987014 |
| EOTAXIN_15 | intron     | RP11-136C24.3 | 3  | 43005217 |
| EOTAXIN_15 | intron     | RP11-136C24.3 | 3  | 43006533 |
| EOTAXIN_15 | intron     | FAM198A       | 3  | 43066572 |
| EOTAXIN_15 | intron     | FAM198A       | 3  | 43068413 |
| EOTAXIN_15 | missense   | FAM198A       | 3  | 43095101 |
| EOTAXIN_15 | downstream | POMGNT2       | 3  | 43116940 |
| EOTAXIN_2  | intron     | CCL26         | 7  | 75408687 |
| EOTAXIN_2  | intron     | RHBDD2        | 7  | 75491565 |
| EOTAXIN_3  | downstream | LZTFL1        | 3  | 45863267 |
| EOTAXIN_3  | intron     | LZTFL1        | 3  | 45910946 |
| EOTAXIN_3  | intron     | LZTFL1        | 3  | 45954352 |
| EOTAXIN_3  | intron     | LZTFL1        | 3  | 45956362 |
| EOTAXIN_3  | upstream   | LZTFL1        | 3  | 45957689 |
| EOTAXIN_3  | intron     | FYCO1         | 3  | 45963497 |
| EOTAXIN_3  | intron     | FYCO1         | 3  | 45976746 |

|             |            |              |    |           |
|-------------|------------|--------------|----|-----------|
| EOTAXIN_3   | intron     | FYCO1        | 3  | 45979877  |
| EOTAXIN_3   | intron     | FYCO1        | 3  | 45980115  |
| EOTAXIN_3   | intron     | FYCO1        | 3  | 45982119  |
| EOTAXIN_3   | intron     | FYCO1        | 3  | 45984300  |
| EOTAXIN_3   | intron     | FYCO1        | 3  | 45993606  |
| EOTAXIN_3   | intron     | FYCO1        | 3  | 45996501  |
| EOTAXIN_3   | intron     | FYCO1        | 3  | 46032847  |
| EOTAXIN_3   | intron     | FYCO1        | 3  | 46034086  |
| EOTAXIN_3   | intergenic | TKT          | 3  | 46050355  |
| EOTAXIN_6   | intron     | SLC25A26     | 3  | 66357919  |
| EOTAXIN_6   | intron     | SLC25A26     | 3  | 66386554  |
| EOTAXIN_6   | intron     | SLC25A26     | 3  | 66398920  |
| EOTAXIN_6   | intron     | SLC25A26     | 3  | 66403649  |
| EOTAXIN_6   | intron     | SLC25A26     | 3  | 66408211  |
| EOTAXIN_6   | intron     | LRIG1        | 3  | 66439115  |
| EOTAXIN_6   | intron     | LRIG1        | 3  | 66447299  |
| EOTAXIN_6   | intron     | LRIG1        | 3  | 66474954  |
| EOTAXIN_6   | intron     | LRIG1        | 3  | 66481294  |
| EOTAXIN_7   | upstream   | DSTNP4       | 3  | 39250894  |
| EOTAXIN_7   | intergenic | DSTNP4       | 3  | 39261447  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39267953  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39281112  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39289218  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39289693  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39291976  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39292424  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39293699  |
| EOTAXIN_7   | downstream | CX3CR1       | 3  | 39302415  |
| EOTAXIN_7   | intron     | CX3CR1       | 3  | 39308205  |
| EOTAXIN_7   | upstream   | CX3CR1       | 3  | 39326283  |
| EOTAXIN_7   | intergenic | RP11-944L7.4 | 3  | 39344438  |
| EOTAXIN_7   | NA         | NA           | NA | NA        |
| EOTAXIN_8   | NA         | NA           | 19 | 37736901  |
| EOTAXIN_8   | NA         | NA           | 19 | 37801242  |
| EOTAXIN_8   | NA         | NA           | 19 | 37803248  |
| EOTAXIN_8   | NA         | NA           | 19 | 37810176  |
| EOTAXIN_8   | NA         | NA           | 19 | 37816230  |
| EOTAXIN_9   | NA         | NA           | 17 | 32552845  |
| EOTAXIN_9   | NA         | NA           | 17 | 32578358  |
| FGF_BASIC_1 | NA         | NA           | 1  | 198882670 |
| FGF_BASIC_1 | intron     | MIR181A1HG   | 1  | 198900138 |
| FGF_BASIC_1 | intergenic | RP11-16L9.1  | 1  | 198926671 |
| FGF_BASIC_1 | intergenic | RP11-16L9.1  | 1  | 198934697 |
| FGF_BASIC_1 | intergenic | RP11-16L9.2  | 1  | 198938170 |
| FGF_BASIC_1 | upstream   | RP11-16L9.2  | 1  | 198951267 |
| FGF_BASIC_1 | downstream | RP11-16L9.2  | 1  | 198956913 |

|               |             |               |    |           |
|---------------|-------------|---------------|----|-----------|
| FGF_BASIC_1   | NA          | NA            | 1  | 198972334 |
| FGF_BASIC_1   | intron      | LINC01222     | 1  | 198979447 |
| FGF_BASIC_1   | intron      | LINC01221     | 1  | 199000350 |
| FGF_BASIC_1   | intron      | LINC01221     | 1  | 199004994 |
| FGF_BASIC_1   | intron      | LINC01221     | 1  | 199007208 |
| FGF_BASIC_1   | intron      | LINC01221     | 1  | 199012816 |
| FGF_BASIC_1   | downstream  | LINC01221     | 1  | 199045870 |
| FGF_BASIC_11  | intergenic  | RP11-170N16.1 | 4  | 122884709 |
| FGF_BASIC_11  | intergenic  | RP11-170N16.1 | 4  | 122911482 |
| FGF_BASIC_11  | intergenic  | RP11-170N16.1 | 4  | 122914769 |
| FGF_BASIC_11  | intergenic  | RP11-170N16.1 | 4  | 122914913 |
| FGF_BASIC_12  | intron      | FGF2          | 4  | 123752017 |
| FGF_BASIC_12  | intron      | FGF2          | 4  | 123764127 |
| FGF_BASIC_2   | intron      | SYNPO2        | 4  | 119970670 |
| FGF_BASIC_2   | 3_prime_UTR | SYNPO2        | 4  | 119981086 |
| FGF_BASIC_2   | 3_prime_UTR | SYNPO2        | 4  | 119981296 |
| FGF_BASIC_2   | 3_prime_UTR | SYNPO2        | 4  | 119981364 |
| FGF_BASIC_2   | intergenic  | RP11-170N16.1 | 4  | 119987701 |
| FGF_BASIC_2   | intergenic  | RP11-170N16.1 | 4  | 120020471 |
| FGF_BASIC_2   | intergenic  | RP11-170N16.1 | 4  | 120048036 |
| FGF_BASIC_2   | intron      | MYOZ2         | 4  | 120076967 |
| FGF_BASIC_2   | intron      | MYOZ2         | 4  | 120096000 |
| FGF_BASIC_9   | intergenic  | RP11-404I7.1  | 4  | 127254177 |
| FGF_BASIC_9   | intergenic  | RP11-404I7.1  | 4  | 127266101 |
| G_CSF_3       | intergenic  | SLC2A2        | 3  | 165706855 |
| G_CSF_3       | intergenic  | SLC2A2        | 3  | 165723243 |
| G_CSF_7       | intron      | MARK4         | 19 | 45700657  |
| G_CSF_7       | intron      | EXOC3L2       | 19 | 45726224  |
| G_CSF_9       | NA          | NA            | 19 | 40417662  |
| G_CSF_9       | NA          | NA            | 19 | 40421612  |
| GROA_CXCL1_10 | NA          | NA            | 4  | 64401593  |
| GROA_CXCL1_10 | intergenic  | DPP3P1        | 4  | 64456551  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74626290  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74667945  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74692819  |
| GROA_CXCL1_15 | intron      | CXCL6         | 4  | 74712514  |
| GROA_CXCL1_15 | upstream    | PF4V1         | 4  | 74715170  |
| GROA_CXCL1_15 | downstream  | PF4V1         | 4  | 74720478  |
| GROA_CXCL1_15 | upstream    | CXCL1         | 4  | 74735108  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74742061  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74745550  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74750147  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74751028  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74759950  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74765488  |
| GROA_CXCL1_15 | intergenic  | RP11-576N17.4 | 4  | 74766693  |

|               |                            |               |    |          |
|---------------|----------------------------|---------------|----|----------|
| GROA_CXCL1_15 | intergenic                 | RP11-576N17.4 | 4  | 74766726 |
| GROA_CXCL1_15 | intergenic                 | RP11-576N17.4 | 4  | 74770221 |
| GROA_CXCL1_15 | intergenic                 | RP11-576N17.4 | 4  | 74773643 |
| GROA_CXCL1_3  | intergenic                 | AC112518.3    | 4  | 68223858 |
| GROA_CXCL1_3  | intergenic                 | AC112518.3    | 4  | 68229448 |
| GROA_CXCL1_3  | non_coding_transcript_exon | RP11-584P21.2 | 4  | 68284668 |
| GROA_CXCL1_3  | downstream                 | CENPC         | 4  | 68332896 |
| GROA_CXCL1_3  | intron                     | CENPC         | 4  | 68363189 |
| GROA_CXCL1_5  | intron                     | SLC4A4        | 4  | 72089575 |
| GROA_CXCL1_5  | intron                     | SLC4A4        | 4  | 72178161 |
| GROA_CXCL1_6  | downstream                 | RNU6-543P     | 10 | 64874754 |
| GROA_CXCL1_6  | upstream                   | RP11-144G16.1 | 10 | 64886103 |
| GROA_CXCL1_6  | upstream                   | NRBF2         | 10 | 64888493 |
| GROA_CXCL1_6  | intron                     | JMJD1C        | 10 | 64961659 |
| HGF_11        | intergenic                 | HGF           | 7  | 81420328 |
| HGF_11        | intergenic                 | HGF           | 7  | 81426413 |
| HGF_11        | intergenic                 | HGF           | 7  | 81430696 |
| HGF_11        | intergenic                 | HGF           | 7  | 81430939 |
| HGF_11        | intergenic                 | HGF           | 7  | 81435605 |
| HGF_11        | intergenic                 | HGF           | 7  | 81437873 |
| HGF_11        | intergenic                 | HGF           | 7  | 81459395 |
| HGF_11        | intergenic                 | HGF           | 7  | 81470110 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81503299 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81515706 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81530304 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81530948 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81531935 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81532442 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81535350 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81537422 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81541336 |
| HGF_11        | intergenic                 | AC006145.1    | 7  | 81561103 |
| HGF_11        | intergenic                 | CACNA2D1      | 7  | 81567479 |
| HGF_2         | intergenic                 | AC098828.2    | 2  | 20314068 |
| HGF_2         | intergenic                 | AC098828.2    | 2  | 20315251 |
| HGF_2         | intergenic                 | AC098828.2    | 2  | 20318517 |
| HGF_2         | intergenic                 | AC098828.2    | 2  | 20321943 |
| HGF_2         | intergenic                 | AC098828.3    | 2  | 20334873 |
| HGF_2         | intergenic                 | AC098828.3    | 2  | 20347893 |
| HGF_2         | downstream                 | AC098828.3    | 2  | 20350751 |
| HGF_2         | downstream                 | AC098828.3    | 2  | 20350783 |
| HGF_2         | intergenic                 | AC098828.3    | 2  | 20363074 |
| HGF_2         | intergenic                 | AC098828.3    | 2  | 20363666 |
| HGF_2         | intergenic                 | RN7SL140P     | 2  | 20367135 |
| HGF_2         | intergenic                 | RN7SL140P     | 2  | 20369562 |
| HGF_2         | intergenic                 | RN7SL140P     | 2  | 20369701 |

|         |             |                |    |           |
|---------|-------------|----------------|----|-----------|
| HGF_2   | upstream    | RN7SL140P      | 2  | 20371380  |
| HGF_2   | upstream    | RN7SL140P      | 2  | 20372409  |
| HGF_2   | upstream    | RN7SL140P      | 2  | 20374249  |
| HGF_2   | intergenic  | SDC1           | 2  | 20393183  |
| HGF_2   | downstream  | SDC1           | 2  | 20396122  |
| HGF_2   | downstream  | SDC1           | 2  | 20399054  |
| HGF_2   | intron      | SDC1           | 2  | 20412735  |
| HGF_4   | NA          | NA             | 12 | 111786892 |
| HGF_4   | NA          | NA             | 12 | 111790511 |
| HGF_4   | NA          | NA             | 12 | 111870790 |
| HGF_4   | NA          | NA             | 12 | 111884608 |
| HGF_4   | NA          | NA             | 12 | 111932800 |
| HGF_4   | NA          | NA             | NA | NA        |
| HGF_6   | NA          | NA             | 11 | 72905849  |
| HGF_6   | NA          | NA             | 11 | 72945341  |
| HGF_8   | intron      | HGF            | 7  | 81339005  |
| HGF_8   | intron      | HGF            | 7  | 81363334  |
| HGF_8   | intron      | HGF            | 7  | 81367652  |
| HGF_8   | intron      | HGF            | 7  | 81375929  |
| HGF_9   | intron      | HTT            | 4  | 3153565   |
| HGF_9   | intron      | HTT            | 4  | 3165947   |
| HGF_9   | intron      | MSANTD1        | 4  | 3256966   |
| HGF_9   | 3_prime_UTR | MSANTD1        | 4  | 3258321   |
| HGF_9   | missense    | MSANTD1        | 4  | 3263138   |
| HGF_9   | 3_prime_UTR | MSANTD1        | 4  | 3272782   |
| HGF_9   | intergenic  | ABLIM2         | 4  | 3282809   |
| HGF_9   | NA          | NA             | 4  | 3283339   |
| HGF_9   | intergenic  | ABLIM2         | 4  | 3284340   |
| HGF_9   | intergenic  | ABLIM2         | 4  | 3285477   |
| HGF_9   | NA          | NA             | 4  | 3334536   |
| IFN_G_5 | upstream    | SLC44A4        | 6  | 31847180  |
| IFN_G_5 | intron      | XXbac-BPG116M5 | 6  | 31916951  |
| IL_10_1 | intergenic  | RP11-344J7.2   | 6  | 43846430  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43850456  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43884367  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43885087  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43888107  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43888210  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43888710  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43900048  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43909383  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43911598  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43918897  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43919162  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43922205  |
| IL_10_1 | intergenic  | ACTG1P9        | 6  | 43925327  |

|          |             |               |    |          |
|----------|-------------|---------------|----|----------|
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43925373 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43925607 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43925778 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43926219 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43926635 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43926638 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43928985 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43930962 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43931347 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43932223 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43941682 |
| IL_10_1  | intergenic  | ACTG1P9       | 6  | 43947398 |
| IL_10_1  | downstream  | RP5-1120P11.1 | 6  | 43961520 |
| IL_10_1  | upstream    | C6orf223      | 6  | 43963995 |
| IL_10_1  | downstream  | C6orf223      | 6  | 43974166 |
| IL_10_1  | downstream  | C6orf223      | 6  | 43977172 |
| IL_10_1  | downstream  | C6orf223      | 6  | 43977535 |
| IL_10_1  | intron      | RP5-1120P11.1 | 6  | 43991377 |
| IL_16_11 | intergenic  | TLE3          | 15 | 70273395 |
| IL_16_11 | NA          | NA            | 15 | 70283382 |
| IL_16_11 | intron      | TLE3          | 15 | 70345281 |
| IL_16_11 | intron      | TLE3          | 15 | 70352198 |
| IL_16_11 | intron      | TLE3          | 15 | 70356560 |
| IL_16_11 | intron      | TLE3          | 15 | 70357918 |
| IL_16_11 | intergenic  | TLE3          | 15 | 70406641 |
| IL_16_11 | intergenic  | TLE3          | 15 | 70431838 |
| IL_16_11 | intergenic  | TLE3          | 15 | 70433706 |
| IL_16_11 | intergenic  | RNU6-745P     | 15 | 70440530 |
| IL_16_13 | intron      | IL16          | 15 | 81518209 |
| IL_16_13 | intron      | IL16          | 15 | 81526302 |
| IL_16_13 | intron      | IL16          | 15 | 81527032 |
| IL_16_13 | intron      | IL16          | 15 | 81529410 |
| IL_16_13 | intron      | IL16          | 15 | 81533695 |
| IL_16_13 | intron      | IL16          | 15 | 81543469 |
| IL_16_13 | intron      | IL16          | 15 | 81550997 |
| IL_16_13 | intron      | IL16          | 15 | 81553532 |
| IL_16_13 | intron      | IL16          | 15 | 81559324 |
| IL_16_13 | intron      | IL16          | 15 | 81561124 |
| IL_16_13 | intron      | IL16          | 15 | 81564437 |
| IL_16_13 | intron      | IL16          | 15 | 81566861 |
| IL_16_13 | intron      | IL16          | 15 | 81574356 |
| IL_16_13 | intron      | IL16          | 15 | 81587757 |
| IL_16_13 | synonymous  | IL16          | 15 | 81592802 |
| IL_16_13 | intron      | IL16          | 15 | 81592891 |
| IL_16_13 | 3_prime_UTR | IL16          | 15 | 81602008 |
| IL_16_13 | intron      | STARD5        | 15 | 81608724 |

|          |            |               |    |          |
|----------|------------|---------------|----|----------|
| IL_16_13 | intron     | STARD5        | 15 | 81612491 |
| IL_16_13 | upstream   | STARD5        | 15 | 81618208 |
| IL_16_13 | upstream   | STARD5        | 15 | 81619705 |
| IL_16_13 | upstream   | STARD5        | 15 | 81620796 |
| IL_16_13 | missense   | TMC3          | 15 | 81625028 |
| IL_16_13 | intron     | TMC3          | 15 | 81645032 |
| IL_16_13 | intron     | TMC3          | 15 | 81655838 |
| IL_16_13 | intron     | TMC3          | 15 | 81657169 |
| IL_16_13 | intron     | TMC3          | 15 | 81657601 |
| IL_16_13 | intron     | TMC3          | 15 | 81662596 |
| IL_16_13 | upstream   | TMC3          | 15 | 81668091 |
| IL_16_13 | upstream   | CTD-2240J17.1 | 15 | 81702577 |
| IL_16_13 | intron     | CTD-2240J17.1 | 15 | 81710618 |
| IL_16_15 | intron     | FGD3          | 9  | 95771569 |
| IL_16_15 | intron     | FGD3          | 9  | 95784509 |
| IL_16_15 | intron     | SUSD3         | 9  | 95827255 |
| IL_16_15 | intron     | SUSD3         | 9  | 95835782 |
| IL_16_15 | upstream   | CARD19        | 9  | 95857720 |
| IL_16_15 | intron     | NINJ1         | 9  | 95887154 |
| IL_16_15 | intron     | NINJ1         | 9  | 95895796 |
| IL_16_15 | upstream   | RP11-370F5.4  | 9  | 95905552 |
| IL_16_15 | upstream   | RP11-370F5.4  | 9  | 95908125 |
| IL_16_15 | intergenic | WNK2          | 9  | 95939766 |
| IL_16_15 | intron     | WNK2          | 9  | 95948978 |
| IL_16_15 | intron     | WNK2          | 9  | 95951065 |
| IL_16_15 | intron     | WNK2          | 9  | 95956632 |
| IL_16_3  | intron     | THSD4         | 15 | 71424046 |
| IL_16_3  | intron     | THSD4         | 15 | 71463600 |
| IL_16_4  | downstream | RP11-38G5.2   | 15 | 80122599 |
| IL_16_4  | intron     | MTHFS         | 15 | 80129813 |
| IL_16_5  | intron     | ACSBG1        | 15 | 78504289 |
| IL_16_5  | intron     | ACSBG1        | 15 | 78512552 |
| IL_16_5  | intron     | ACSBG1        | 15 | 78525008 |
| IL_16_5  | intron     | ACSBG1        | 15 | 78531287 |
| IL_16_5  | intron     | WDR61         | 15 | 78583531 |
| IL_16_5  | intron     | WDR61         | 15 | 78586915 |
| IL_16_5  | intron     | WDR61         | 15 | 78590086 |
| IL_16_5  | intergenic | RP11-5O23.2   | 15 | 78622728 |
| IL_16_5  | upstream   | CRABP1        | 15 | 78628257 |
| IL_16_6  | intron     | AKAP13        | 15 | 86098507 |
| IL_16_6  | intron     | AKAP13        | 15 | 86104437 |
| IL_16_6  | missense   | AKAP13        | 15 | 86123809 |
| IL_16_6  | intron     | AKAP13        | 15 | 86160441 |
| IL_16_6  | intron     | AKAP13        | 15 | 86167257 |
| IL_16_6  | intron     | AKAP13        | 15 | 86167428 |
| IL_16_6  | intron     | AKAP13        | 15 | 86168731 |



|          |             |               |    |           |
|----------|-------------|---------------|----|-----------|
| IL_16_6  | intron      | AKAP13        | 15 | 86194087  |
| IL_16_6  | intron      | AKAP13        | 15 | 86194281  |
| IL_16_6  | intron      | AKAP13        | 15 | 86199049  |
| IL_16_6  | intron      | AKAP13        | 15 | 86217704  |
| IL_16_6  | intron      | AKAP13        | 15 | 86229273  |
| IL_16_6  | intron      | AKAP13        | 15 | 86258275  |
| IL_16_7  | intron      | CFAP161       | 15 | 81416203  |
| IL_16_7  | intron      | IL16          | 15 | 81492898  |
| IL_16_7  | intron      | IL16          | 15 | 81508398  |
| IL_16_7  | intron      | IL16          | 15 | 81522543  |
| IL_16_7  | intron      | IL16          | 15 | 81524507  |
| IL_16_9  | intron      | CPEB1         | 15 | 83304977  |
| IL_16_9  | intron      | CPEB1         | 15 | 83309945  |
| IL_16_9  | intron      | AP3B2         | 15 | 83338505  |
| IL_16_9  | downstream  | ACTG1P17      | 15 | 83391441  |
| IL_16_9  | downstream  | ACTG1P17      | 15 | 83393723  |
| IL_16_9  | intron      | RP11-752G15.8 | 15 | 83416452  |
| IL_16_9  | 3_prime_UTR | FSD2          | 15 | 83426185  |
| IL_16_9  | intron      | FSD2          | 15 | 83431905  |
| IL_18_1  | intergenic  | JMY           | 5  | 68100579  |
| IL_18_1  | intergenic  | JMY           | 5  | 68120495  |
| IL_18_1  | downstream  | CTC-340D7.1   | 5  | 68124912  |
| IL_18_1  | downstream  | CTC-340D7.1   | 5  | 68126696  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68142407  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68143193  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68186028  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68214983  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68216562  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68224163  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68231686  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68233743  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68236252  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68240482  |
| IL_18_1  | intron      | CTC-340D7.1   | 5  | 68242147  |
| IL_18_10 | upstream    | IL18          | 11 | 112037014 |
| IL_18_10 | intron      | BCO2          | 11 | 112056400 |
| IL_18_4  | upstream    | MTCO1P54      | 2  | 203489457 |
| IL_18_4  | upstream    | FAM117B       | 2  | 203495058 |
| IL_18_5  | NA          | NA            | 8  | 144475702 |
| IL_18_5  | intergenic  | RP11-909N17.2 | 8  | 144484355 |
| IL_18_5  | NA          | NA            | 8  | 144502917 |
| IL_18_5  | NA          | NA            | 8  | 144518792 |
| IL_18_5  | NA          | NA            | 8  | 144554746 |
| IL_18_5  | intron      | ZC3H3         | 8  | 144561183 |
| IL_18_5  | intron      | ZC3H3         | 8  | 144601816 |
| IL_18_5  | NA          | NA            | 8  | 144627241 |

|         |            |            |    |           |
|---------|------------|------------|----|-----------|
| IL_18_5 | NA         | NA         | 8  | 144630847 |
| IL_18_5 | intron     | GSDMD      | 8  | 144636596 |
| IL_18_5 | synonymous | GSDMD      | 8  | 144642854 |
| IL_18_5 | missense   | MROH6      | 8  | 144654249 |
| IL_18_5 | missense   | MROH6      | 8  | 144654594 |
| IL_18_5 | upstream   | MROH6      | 8  | 144656187 |
| IL_18_5 | NA         | NA         | 8  | 144656765 |
| IL_18_6 | NA         | NA         | 11 | 104929590 |
| IL_18_6 | NA         | NA         | NA | NA        |
| IL_18_7 | intergenic | MIR148A    | 7  | 26038458  |
| IL_18_7 | intergenic | MIR148A    | 7  | 26042998  |
| IL_18_7 | intergenic | AC004520.1 | 7  | 26048012  |
| IL_18_7 | intergenic | AC004520.1 | 7  | 26049848  |
| IL_18_7 | intergenic | AC004520.1 | 7  | 26072772  |
| IL_18_7 | intergenic | AC004520.1 | 7  | 26094886  |
| IL_18_7 | intergenic | AC004520.1 | 7  | 26109740  |
| IL_18_7 | intergenic | AC004520.1 | 7  | 26126848  |
| IL_18_7 | intergenic | AC004520.1 | 7  | 26128382  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26146633  |
| IL_18_7 | NA         | NA         | 7  | 26147439  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26148201  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26149244  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26149563  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26152232  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26153279  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26153556  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26157954  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26158521  |
| IL_18_7 | intergenic | NFE2L3     | 7  | 26170930  |
| IL_18_7 | intron     | NFE2L3     | 7  | 26211703  |
| IL_18_7 | intron     | HNRNPA2B1  | 7  | 26232396  |
| IL_18_9 | intergenic | XDH        | 2  | 31543080  |
| IL_18_9 | intergenic | XDH        | 2  | 31546171  |
| IL_18_9 | intron     | XDH        | 2  | 31563398  |
| IL_18_9 | intron     | XDH        | 2  | 31570626  |
| IL_18_9 | intron     | XDH        | 2  | 31572732  |
| IL_18_9 | intron     | XDH        | 2  | 31588577  |
| IL_18_9 | intron     | XDH        | 2  | 31599699  |
| IL_18_9 | intron     | XDH        | 2  | 31609961  |
| IL_18_9 | intron     | XDH        | 2  | 31611367  |
| IL_18_9 | intron     | XDH        | 2  | 31633014  |
| IL_18_9 | intron     | XDH        | 2  | 31633965  |
| IL_18_9 | intron     | XDH        | 2  | 31636915  |
| IL_18_9 | intergenic | XDH        | 2  | 31649666  |
| IL_18_9 | intergenic | XDH        | 2  | 31656380  |
| IL_18_9 | intergenic | XDH        | 2  | 31659226  |

|          |                            |              |    |           |
|----------|----------------------------|--------------|----|-----------|
| IL_18_9  | intergenic                 | XDH          | 2  | 31663470  |
| IL_18_9  | intergenic                 | XDH          | 2  | 31667530  |
| IL_18_9  | intergenic                 | XDH          | 2  | 31675248  |
| IL_18_9  | intergenic                 | XDH          | 2  | 31676320  |
| IL_1B_1  | NA                         | NA           | 2  | 3527789   |
| IL_1B_1  | intron                     | RP13-512J5.1 | 2  | 3587281   |
| IL_1B_1  | intron                     | RP13-512J5.1 | 2  | 3589803   |
| IL_1B_1  | NA                         | NA           | 2  | 3614960   |
| IL_1B_1  | upstream                   | RPS7         | 2  | 3618008   |
| IL_1B_1  | intron                     | RPS7         | 2  | 3628383   |
| IL_1B_1  | downstream                 | RPS7         | 2  | 3630701   |
| IL_1B_1  | intergenic                 | RPS7         | 2  | 3634043   |
| IL_1B_1  | intergenic                 | RPS7         | 2  | 3634606   |
| IL_1B_1  | intergenic                 | COLEC11      | 2  | 3636478   |
| IL_1B_1  | upstream                   | COLEC11      | 2  | 3640142   |
| IL_1B_1  | upstream                   | COLEC11      | 2  | 3642361   |
| IL_1B_1  | intron                     | COLEC11      | 2  | 3648396   |
| IL_1B_1  | intron                     | COLEC11      | 2  | 3659579   |
| IL_1B_1  | NA                         | NA           | 2  | 3660416   |
| IL_1B_1  | NA                         | NA           | 2  | 3667871   |
| IL_1B_1  | NA                         | NA           | 2  | 3680302   |
| IL_1B_1  | intron                     | COLEC11      | 2  | 3683651   |
| IL_1B_1  | NA                         | NA           | 2  | 3699278   |
| IL_1B_1  | NA                         | NA           | 2  | 3705001   |
| IL_1B_1  | NA                         | NA           | 2  | 3713658   |
| IL_1B_1  | NA                         | NA           | NA | NA        |
| IL_1B_3  | NA                         | NA           | 2  | 1075456   |
| IL_1B_3  | intron                     | SNTG2        | 2  | 1099624   |
| IL_1B_3  | NA                         | NA           | 2  | 1116730   |
| IL_1B_3  | NA                         | NA           | 2  | 1124479   |
| IL_1B_3  | intron                     | SNTG2        | 2  | 1147224   |
| IL_1B_3  | NA                         | NA           | 2  | 1152921   |
| IL_1B_3  | intron                     | SNTG2        | 2  | 1156484   |
| IL_1B_3  | NA                         | NA           | 2  | 1156587   |
| IL_1B_3  | NA                         | NA           | 2  | 1157438   |
| IL_1B_3  | intron                     | SNTG2        | 2  | 1199248   |
| IL_1B_3  | NA                         | NA           | 2  | 1232471   |
| IL_1B_3  | NA                         | NA           | 2  | 1256951   |
| IL_1B_3  | NA                         | NA           | NA | NA        |
| IL_1B_5  | non_coding_transcript_exon | C3P1         | 19 | 10148713  |
| IL_1B_5  | synonymous                 | EIF3G        | 19 | 10227847  |
| IL_1RA_1 | intergenic                 | IGKV1OR2-108 | 2  | 114155991 |
| IL_1RA_1 | intergenic                 | IGKV1OR2-108 | 2  | 114156855 |
| IL_1RA_2 | upstream                   | MIR518E      | 19 | 54232808  |
| IL_1RA_2 | upstream                   | MIR521-1     | 19 | 54248085  |
| IL_1RA_2 | upstream                   | RNU6-751P    | 19 | 54257748  |

|          |             |              |    |           |
|----------|-------------|--------------|----|-----------|
| IL_1RA_2 | upstream    | AC011453.1   | 19 | 54259376  |
| IL_1RA_2 | upstream    | SEPT7P8      | 19 | 54277963  |
| IL_1RA_2 | upstream    | CTD-3022G6.1 | 19 | 54291427  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54301284  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54302872  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54316051  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54318322  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54319624  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54320636  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54320716  |
| IL_1RA_2 | intron      | NLRP12       | 19 | 54324916  |
| IL_1RA_2 | upstream    | NLRP12       | 19 | 54327869  |
| IL_1RA_2 | intergenic  | NLRP12       | 19 | 54337509  |
| IL_1RA_2 | intergenic  | NLRP12       | 19 | 54338067  |
| IL_1RA_2 | intergenic  | NLRP12       | 19 | 54338682  |
| IL_1RA_2 | intergenic  | AC008753.6   | 19 | 54345604  |
| IL_1RA_2 | intron      | PRKCG        | 19 | 54394213  |
| IL_1RA_4 | intron      | IL36RN       | 2  | 113818727 |
| IL_1RA_4 | -           | -            | 2  | 113819451 |
| IL_1RA_8 | intron      | FTO          | 16 | 53799847  |
| IL_1RA_8 | intron      | FTO          | 16 | 53830452  |
| IL_2_1   | intron      | KLKB1        | 4  | 187130198 |
| IL_2_1   | intron      | KLKB1        | 4  | 187160394 |
| IL_2_2   | intergenic  | MLXIPL       | 7  | 73050599  |
| IL_2_2   | intron      | STX1A        | 7  | 73128525  |
| IL_2_3   | intron      | LMAN2        | 5  | 176766177 |
| IL_2_3   | upstream    | RGS14        | 5  | 176784612 |
| IL_2_3   | synonymous  | SLC34A1      | 5  | 176815124 |
| IL_2_3   | intron      | SLC34A1      | 5  | 176815766 |
| IL_2_3   | NA          | NA           | 5  | 176830627 |
| IL_2_3   | 5_prime_UTR | F12          | 5  | 176836532 |
| IL_2_3   | intron      | GRK6         | 5  | 176837578 |
| IL_2_3   | intron      | GRK6         | 5  | 176842474 |
| IL_2_3   | intron      | GRK6         | 5  | 176867276 |
| IL_2_4   | intron      | TRIM54       | 2  | 27527321  |
| IL_2_4   | intron      | SNX17        | 2  | 27598097  |
| IL_2RA_2 | intron      | FBXO18       | 10 | 5933150   |
| IL_2RA_2 | intron      | IL15RA       | 10 | 5999657   |
| IL_2RA_2 | intergenic  | IL15RA       | 10 | 6035457   |
| IL_2RA_2 | intergenic  | IL2RA        | 10 | 6042034   |
| IL_2RA_2 | intron      | IL2RA        | 10 | 6058735   |
| IL_2RA_2 | intron      | IL2RA        | 10 | 6069893   |
| IL_2RA_2 | intron      | IL2RA        | 10 | 6070273   |
| IL_2RA_2 | intron      | IL2RA        | 10 | 6071379   |
| IL_2RA_2 | intron      | IL2RA        | 10 | 6075831   |
| IL_2RA_2 | intron      | IL2RA        | 10 | 6079017   |

|          |            |               |    |           |
|----------|------------|---------------|----|-----------|
| IL_2RA_2 | intron     | IL2RA         | 10 | 6090322   |
| IL_2RA_2 | intron     | IL2RA         | 10 | 6095928   |
| IL_4_2   | NA         | NA            | 16 | 17592122  |
| IL_4_2   | NA         | NA            | 16 | 17617489  |
| IL_6_1   | NA         | NA            | 1  | 154400799 |
| IL_6_1   | NA         | NA            | 1  | 154414086 |
| IL_6_1   | NA         | NA            | 1  | 154415777 |
| IL_6_1   | NA         | NA            | 1  | 154416969 |
| IL_6_1   | NA         | NA            | 1  | 154417044 |
| IL_6_1   | NA         | NA            | 1  | 154425456 |
| IL_6_1   | NA         | NA            | 1  | 154487926 |
| IL_6_1   | NA         | NA            | NA | NA        |
| IL_8_1   | intron     | PIGL          | 17 | 16190418  |
| IL_8_1   | intron     | PIGL          | 17 | 16201941  |
| IL_8_2   | intron     | AIM2          | 1  | 159088541 |
| IL_8_2   | upstream   | AIM2          | 1  | 159114670 |
| IL_8_2   | intergenic | CADM3         | 1  | 159131926 |
| IL_8_2   | intron     | CADM3         | 1  | 159150246 |
| IL_8_2   | intron     | CADM3         | 1  | 159169463 |
| IL_8_2   | missense   | ACKR1         | 1  | 159175354 |
| IL_8_2   | downstream | CADM3         | 1  | 159176490 |
| IL_8_2   | downstream | CADM3         | 1  | 159177748 |
| IL_8_2   | intergenic | MPTX1         | 1  | 159230865 |
| IL_8_2   | intron     | FCER1A        | 1  | 159269795 |
| IL_8_2   | intron     | FCER1A        | 1  | 159271482 |
| IL_8_2   | intron     | FCER1A        | 1  | 159273372 |
| IL_8_8   | intergenic | PSMD3         | 17 | 38160754  |
| IL_8_8   | intergenic | RP11-387H17.6 | 17 | 38163295  |
| IL_9_1   | intergenic | SMUG1         | 12 | 54602032  |
| IL_9_1   | intron     | CBX5          | 12 | 54635846  |
| IL_9_1   | intron     | CBX5          | 12 | 54636783  |
| IL_9_1   | intron     | CBX5          | 12 | 54664097  |
| IL_9_1   | intron     | CBX5          | 12 | 54668471  |
| IL_9_1   | intron     | HNRNPA1       | 12 | 54677055  |
| IL_9_1   | intron     | HNRNPA1       | 12 | 54677980  |
| IL_9_1   | downstream | NFE2          | 12 | 54683639  |
| IL_9_1   | downstream | NFE2          | 12 | 54684768  |
| IL_9_1   | downstream | NFE2          | 12 | 54684955  |
| IL_9_1   | intron     | NFE2          | 12 | 54687232  |
| IL_9_1   | intron     | NFE2          | 12 | 54687489  |
| IL_9_1   | intron     | NFE2          | 12 | 54692061  |
| IL_9_1   | intron     | COPZ1         | 12 | 54706415  |
| IL_9_1   | intron     | COPZ1         | 12 | 54712308  |
| IL_9_1   | intron     | COPZ1         | 12 | 54712700  |
| IL_9_1   | intron     | COPZ1         | 12 | 54715745  |
| IL_9_1   | intron     | COPZ1         | 12 | 54716092  |

|          |            |               |    |           |
|----------|------------|---------------|----|-----------|
| IL_9_1   | intron     | COPZ1         | 12 | 54719689  |
| IL_9_1   | intron     | COPZ1         | 12 | 54722983  |
| IL_9_1   | intron     | COPZ1         | 12 | 54723028  |
| IL_9_1   | intron     | COPZ1         | 12 | 54726475  |
| IL_9_1   | intron     | COPZ1         | 12 | 54729872  |
| IL_9_1   | intron     | COPZ1         | 12 | 54730754  |
| IL_9_1   | intron     | COPZ1         | 12 | 54736470  |
| IL_9_1   | intron     | RP11-753H16.3 | 12 | 54750706  |
| IL_9_1   | upstream   | GPR84         | 12 | 54758960  |
| IL_9_1   | upstream   | GPR84         | 12 | 54762150  |
| IL_9_1   | intron     | ZNF385A       | 12 | 54765785  |
| IL_9_3   | downstream | AC009014.3    | 5  | 135525521 |
| IL_9_3   | intergenic | CTB-1I21.1    | 5  | 135534558 |
| IL_9_4   | upstream   | HBS1L         | 6  | 135426558 |
| IL_9_4   | upstream   | HBS1L         | 6  | 135427817 |
| IP_10_1  | intron     | ART3          | 4  | 76937278  |
| IP_10_1  | -          | -             | 4  | 76952288  |
| M_CSF_10 | intron     | VMP1          | 17 | 57850251  |
| M_CSF_10 | intron     | VMP1          | 17 | 57879610  |
| M_CSF_12 | intron     | ST3GAL4       | 11 | 126242546 |
| M_CSF_12 | intron     | ST3GAL4       | 11 | 126243709 |
| M_CSF_14 | intron     | UBE2O         | 17 | 74412284  |
| M_CSF_14 | intron     | UBE2O         | 17 | 74412818  |
| M_CSF_14 | intron     | UBE2O         | 17 | 74412826  |
| M_CSF_14 | intron     | UBE2O         | 17 | 74442440  |
| M_CSF_14 | intron     | RHBDF2        | 17 | 74480715  |
| M_CSF_14 | intron     | RHBDF2        | 17 | 74482126  |
| M_CSF_14 | intron     | RHBDF2        | 17 | 74483143  |
| M_CSF_14 | intron     | RHBDF2        | 17 | 74485549  |
| M_CSF_14 | intron     | ST6GALNAC2    | 17 | 74579983  |
| M_CSF_16 | intron     | ABO           | 9  | 136132957 |
| M_CSF_16 | upstream   | ABO           | 9  | 136155444 |
| M_CSF_18 | intergenic | RP11-374I15.1 | 6  | 95921635  |
| M_CSF_18 | downstream | MANEA-AS1     | 6  | 96018379  |
| M_CSF_19 | intron     | GLB1L2        | 11 | 134216786 |
| M_CSF_19 | intron     | GLB1L2        | 11 | 134234835 |
| M_CSF_19 | downstream | GLB1L2        | 11 | 134247728 |
| M_CSF_19 | intron     | B3GAT1        | 11 | 134275746 |
| M_CSF_19 | intron     | B3GAT1        | 11 | 134280067 |
| M_CSF_19 | -          | -             | 11 | 134280504 |
| M_CSF_19 | intron     | B3GAT1        | 11 | 134281332 |
| M_CSF_2  | intron     | PNPLA5        | 22 | 44281144  |
| M_CSF_2  | intron     | PNPLA3        | 22 | 44332493  |
| M_CSF_2  | intron     | PNPLA3        | 22 | 44334486  |
| M_CSF_2  | intron     | PNPLA3        | 22 | 44334529  |
| M_CSF_2  | intron     | PNPLA3        | 22 | 44335331  |

|          |            |              |    |           |
|----------|------------|--------------|----|-----------|
| M_CSF_2  | NA         | NA           | 22 | 44337533  |
| M_CSF_2  | intron     | PNPLA3       | 22 | 44338105  |
| M_CSF_2  | downstream | PNPLA3       | 22 | 44345771  |
| M_CSF_2  | intron     | SAMM50       | 22 | 44355150  |
| M_CSF_2  | intron     | SAMM50       | 22 | 44357194  |
| M_CSF_2  | intron     | SAMM50       | 22 | 44358290  |
| M_CSF_2  | NA         | NA           | 22 | 44360797  |
| M_CSF_2  | intron     | SAMM50       | 22 | 44366874  |
| M_CSF_2  | intron     | SAMM50       | 22 | 44374267  |
| M_CSF_2  | intron     | SAMM50       | 22 | 44374908  |
| M_CSF_2  | intron     | SAMM50       | 22 | 44390568  |
| M_CSF_2  | intron     | PARVB        | 22 | 44398951  |
| M_CSF_20 | NA         | NA           | 1  | 9118423   |
| M_CSF_20 | NA         | NA           | 1  | 9118596   |
| M_CSF_20 | NA         | NA           | 1  | 9124705   |
| M_CSF_20 | intron     | MIR34AHG     | 1  | 9216615   |
| M_CSF_20 | intron     | MIR34AHG     | 1  | 9234143   |
| M_CSF_20 | intron     | MIR34AHG     | 1  | 9241278   |
| M_CSF_20 | NA         | NA           | 1  | 9241839   |
| M_CSF_20 | intron     | LINC01759    | 1  | 9244795   |
| M_CSF_20 | intron     | LINC01759    | 1  | 9251279   |
| M_CSF_20 | intergenic | RP3-510D11.4 | 1  | 9266901   |
| M_CSF_20 | intergenic | RP3-510D11.4 | 1  | 9272378   |
| M_CSF_20 | NA         | NA           | 1  | 9275926   |
| M_CSF_20 | intergenic | H6PD         | 1  | 9284375   |
| M_CSF_20 | intergenic | H6PD         | 1  | 9287049   |
| M_CSF_20 | intergenic | H6PD         | 1  | 9288466   |
| M_CSF_20 | upstream   | H6PD         | 1  | 9291036   |
| M_CSF_20 | NA         | NA           | 1  | 9296479   |
| M_CSF_20 | NA         | NA           | NA | NA        |
| M_CSF_21 | intron     | HMGXB3       | 5  | 149394658 |
| M_CSF_21 | intron     | CSF1R        | 5  | 149475904 |
| M_CSF_21 | intron     | CSF1R        | 5  | 149482262 |
| M_CSF_21 | intron     | CSF1R        | 5  | 149492547 |
| M_CSF_21 | intron     | PDGFRB       | 5  | 149498151 |
| M_CSF_21 | intron     | PDGFRB       | 5  | 149533976 |
| M_CSF_22 | synonymous | NTN5         | 19 | 49164952  |
| M_CSF_22 | missense   | NTN5         | 19 | 49165157  |
| M_CSF_22 | missense   | NTN5         | 19 | 49167860  |
| M_CSF_22 | intron     | NTN5         | 19 | 49168942  |
| M_CSF_22 | upstream   | NTN5         | 19 | 49178752  |
| M_CSF_22 | intron     | FUT2         | 19 | 49199934  |
| M_CSF_22 | synonymous | FUT2         | 19 | 49206417  |
| M_CSF_22 | stop_lost  | FUT2         | 19 | 49206674  |
| M_CSF_22 | -          | -            | 19 | 49207257  |
| M_CSF_22 | downstream | FUT2         | 19 | 49209226  |

|          |                            |           |    |           |
|----------|----------------------------|-----------|----|-----------|
| M_CSF_22 | downstream                 | FUT2      | 19 | 49209318  |
| M_CSF_22 | downstream                 | FUT2      | 19 | 49209464  |
| M_CSF_22 | intron                     | MAMSTR    | 19 | 49219459  |
| M_CSF_22 | upstream                   | MAMSTR    | 19 | 49223633  |
| M_CSF_22 | intron                     | IZUMO1    | 19 | 49246487  |
| M_CSF_22 | intron                     | BCAT2     | 19 | 49303740  |
| M_CSF_23 | intron                     | TP53      | 17 | 7577708   |
| M_CSF_23 | intron                     | DNAH2     | 17 | 7624799   |
| M_CSF_24 | intron                     | ST3GAL6   | 3  | 98480330  |
| M_CSF_24 | intron                     | ST3GAL6   | 3  | 98486551  |
| M_CSF_24 | intron                     | ST3GAL6   | 3  | 98493447  |
| M_CSF_24 | intron                     | DCBLD2    | 3  | 98543301  |
| M_CSF_24 | intron                     | DCBLD2    | 3  | 98563651  |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134927245 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134949239 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134959755 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134966562 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134971030 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134973524 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134974074 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134975144 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134981216 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134985336 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134987739 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134988291 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134988860 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 134995611 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135006732 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135017417 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135018976 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135031863 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135038848 |
| M_CSF_25 | -                          | -         | 2  | 135040472 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135042375 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135054627 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135056853 |
| M_CSF_25 | intron                     | MGAT5     | 2  | 135066476 |
| M_CSF_3  | intron                     | CSF1      | 1  | 110462488 |
| M_CSF_3  | intergenic                 | AHCYL1    | 1  | 110501597 |
| M_CSF_4  | non_coding_transcript_exon | LINC01768 | 1  | 110371748 |
| M_CSF_4  | missense                   | CSF1      | 1  | 110466466 |
| M_CSF_4  | 3_prime_UTR                | CSF1      | 1  | 110472956 |
| M_CSF_4  | intergenic                 | CSF1      | 1  | 110497808 |
| M_CSF_4  | intergenic                 | CSF1      | 1  | 110499847 |
| M_CSF_5  | intron                     | B4GALT1   | 9  | 33117965  |
| M_CSF_5  | intron                     | B4GALT1   | 9  | 33123013  |



|         |             |              |    |           |
|---------|-------------|--------------|----|-----------|
| MCP_3_2 | upstream    | CCL2         | 17 | 32579958  |
| MCP_3_2 | intergenic  | CCL7         | 17 | 32591916  |
| MCP_3_2 | upstream    | CCL7         | 17 | 32592905  |
| MCP_3_2 | upstream    | CCL7         | 17 | 32594257  |
| MCP_3_2 | intron      | CCL11        | 17 | 32614442  |
| MCP_3_2 | missense    | CCL8         | 17 | 32647357  |
| MCP_3_2 | intergenic  | CCL13        | 17 | 32676323  |
| MCP_3_2 | intergenic  | CCL13        | 17 | 32677148  |
| MCP_3_2 | intergenic  | AC011193.1   | 17 | 32762095  |
| MCP_3_2 | intergenic  | AC011193.1   | 17 | 32767379  |
| MCP_3_5 | intron      | LRRC37B      | 17 | 30364610  |
| MCP_3_5 | upstream    | CTD-2095E4.5 | 17 | 30454415  |
| MCP_3_8 | intergenic  | LINC01989    | 17 | 32534983  |
| MCP_3_8 | intergenic  | LINC01989    | 17 | 32535173  |
| MIF_1   | downstream  | GSTT2B       | 22 | 24297130  |
| MIF_1   | intron      | DDTL         | 22 | 24312204  |
| MIG_1   | upstream    | AC110615.1   | 4  | 76870291  |
| MIG_1   | intron      | CXCL9        | 4  | 76928428  |
| MIG_5   | intron      | MAGI3        | 1  | 114204593 |
| MIG_5   | intron      | MAGI3        | 1  | 114210745 |
| MIG_5   | -           | MAGI3        | 1  | 114221123 |
| MIG_5   | intron      | PHTF1        | 1  | 114265244 |
| MIG_5   | intron      | PHTF1        | 1  | 114299516 |
| MIG_5   | 5_prime_UTR | PHTF1        | 1  | 114301335 |
| MIG_5   | upstream    | PHTF1        | 1  | 114303808 |
| MIG_5   | intron      | RSBN1        | 1  | 114329408 |
| MIG_5   | intron      | PTPN22       | 1  | 114357750 |
| MIG_5   | intron      | PTPN22       | 1  | 114400804 |
| MIG_7   | intron      | LPA          | 6  | 161072331 |
| MIG_7   | intron      | LPA          | 6  | 161083531 |
| MIG_7   | intron      | RP1-81D8.4   | 6  | 161092438 |
| MIG_7   | intron      | PLG          | 6  | 161127125 |
| MIG_7   | intron      | PLG          | 6  | 161131516 |
| MIG_7   | intron      | PLG          | 6  | 161131818 |
| MIG_7   | intron      | PLG          | 6  | 161133244 |
| MIG_7   | intron      | PLG          | 6  | 161133544 |
| MIG_7   | intron      | PLG          | 6  | 161153527 |
| MIG_7   | intron      | PLG          | 6  | 161153561 |
| MIG_7   | intron      | PLG          | 6  | 161156667 |
| MIG_7   | intron      | PLG          | 6  | 161159016 |
| MIG_7   | intron      | PLG          | 6  | 161163777 |
| MIG_7   | upstream    | RP1-81D8.5   | 6  | 161183455 |
| MIG_7   | intergenic  | RP1-155D22.1 | 6  | 161188103 |
| MIG_7   | intron      | RP1-81D8.6   | 6  | 161196162 |
| MIG_7   | intron      | RP1-81D8.6   | 6  | 161196476 |
| MIG_7   | intron      | RP1-81D8.6   | 6  | 161196562 |

|           |             |               |    |           |
|-----------|-------------|---------------|----|-----------|
| MIG_7     | intron      | RP1-81D8.6    | 6  | 161201960 |
| MIP_1A_3  | intergenic  | RP11-500G9.1  | 4  | 38724573  |
| MIP_1A_3  | NA          | NA            | 4  | 38726498  |
| MIP_1A_3  | downstream  | RNA5SP158     | 4  | 38757400  |
| MIP_1A_3  | 3_prime_UTR | TLR10         | 4  | 38774337  |
| MIP_1A_3  | synonymous  | TLR10         | 4  | 38775040  |
| MIP_1A_3  | missense    | TLR10         | 4  | 38776107  |
| MIP_1A_3  | upstream    | TLR10         | 4  | 38788234  |
| MIP_1A_3  | upstream    | TLR10         | 4  | 38789524  |
| MIP_1A_3  | downstream  | TLR1          | 4  | 38789740  |
| MIP_1A_3  | missense    | TLR1          | 4  | 38799710  |
| MIP_1A_3  | intron      | TLR1          | 4  | 38803784  |
| MIP_1A_3  | intron      | TLR1          | 4  | 38807654  |
| MIP_1A_3  | intron      | TLR1          | 4  | 38807911  |
| MIP_1A_3  | intron      | TLR1          | 4  | 38808216  |
| MIP_1A_3  | intron      | TLR1          | 4  | 38808671  |
| MIP_1A_3  | intron      | TLR1          | 4  | 38809598  |
| MIP_1A_3  | intron      | TLR6          | 4  | 38846719  |
| MIP_1A_3  | intron      | TLR6          | 4  | 38854183  |
| MIP_1A_3  | intergenic  | RP11-500G9.1  | 4  | 38864090  |
| MIP_1A_3  | intron      | FAM114A1      | 4  | 38879088  |
| MIP_1A_5  | intergenic  | AC013480.2    | 2  | 42233374  |
| MIP_1A_5  | intergenic  | AC013480.2    | 2  | 42263721  |
| MIP_1A_5  | upstream    | PKDCC         | 2  | 42272662  |
| MIP_1A_5  | synonymous  | PKDCC         | 2  | 42275819  |
| MIP_1A_5  | intergenic  | PKDCC         | 2  | 42318083  |
| MIP_1A_5  | intergenic  | PKDCC         | 2  | 42324419  |
| MIP_1A_5  | intergenic  | AC083949.1    | 2  | 42329711  |
| MIP_1A_5  | intergenic  | AC083949.1    | 2  | 42335185  |
| MIP_1A_5  | intron      | AC083949.1    | 2  | 42381977  |
| MIP_1A_5  | intron      | EML4          | 2  | 42402827  |
| MIP_1A_6  | upstream    | CCL23         | 17 | 34346267  |
| MIP_1A_6  | upstream    | CCL18         | 17 | 34389361  |
| MIP_1A_6  | intergenic  | CCL4          | 17 | 34425574  |
| MIP_1A_6  | upstream    | CCL4          | 17 | 34429459  |
| MIP_1A_6  | upstream    | CTB-91J4.1    | 17 | 34440683  |
| MIP_1A_6  | intron      | CTB-91J4.1    | 17 | 34461538  |
| MIP_1B_11 | upstream    | LYZL6         | 17 | 34273248  |
| MIP_1B_11 | intergenic  | CTB-186H2.2   | 17 | 34284482  |
| MIP_1B_11 | downstream  | CTB-186H2.2   | 17 | 34287702  |
| MIP_1B_11 | intron      | AC069363.1    | 17 | 34408118  |
| MIP_1B_6  | intergenic  | RP11-500B12.1 | 9  | 120378483 |
| MIP_1B_6  | intergenic  | RP11-500B12.1 | 9  | 120385067 |
| MIP_1B_6  | intron      | RP11-500B12.1 | 9  | 120413739 |
| MIP_1B_6  | intergenic  | RP11-500B12.1 | 9  | 120430497 |
| MIP_1B_6  | intergenic  | RPL35AP22     | 9  | 120437061 |

|            |             |               |    |           |
|------------|-------------|---------------|----|-----------|
| MIP_1B_6   | downstream  | RPL35AP22     | 9  | 120445534 |
| MIP_1B_6   | upstream    | RPL35AP22     | 9  | 120449607 |
| MIP_1B_6   | intergenic  | RPL35AP22     | 9  | 120453971 |
| MIP_1B_6   | upstream    | TLR4          | 9  | 120464136 |
| MIP_1B_6   | intron      | TLR4          | 9  | 120467576 |
| MIP_1B_6   | missense    | TLR4          | 9  | 120475602 |
| MIP_1B_6   | intergenic  | RP11-281A20.2 | 9  | 120535975 |
| PDGF_BB_1  | intron      | AK3           | 9  | 4715341   |
| PDGF_BB_1  | intron      | AK3           | 9  | 4724607   |
| PDGF_BB_1  | intron      | AK3           | 9  | 4732225   |
| PDGF_BB_1  | downstream  | RP11-307I14.2 | 9  | 4780007   |
| PDGF_BB_1  | intron      | RCL1          | 9  | 4796624   |
| PDGF_BB_1  | intron      | RCL1          | 9  | 4827637   |
| PDGF_BB_1  | intron      | RCL1          | 9  | 4827719   |
| PDGF_BB_1  | intron      | RCL1          | 9  | 4857662   |
| PDGF_BB_11 | intron      | CD36          | 7  | 80154426  |
| PDGF_BB_11 | intron      | CD36          | 7  | 80224314  |
| PDGF_BB_11 | intron      | CD36          | 7  | 80249671  |
| PDGF_BB_12 | intron      | SLC35D2       | 9  | 99085796  |
| PDGF_BB_12 | intron      | SLC35D2       | 9  | 99091009  |
| PDGF_BB_12 | intron      | SLC35D2       | 9  | 99105420  |
| PDGF_BB_12 | intron      | SLC35D2       | 9  | 99118445  |
| PDGF_BB_12 | intron      | ZNF367        | 9  | 99171381  |
| PDGF_BB_14 | intron      | MFSD13A       | 10 | 104234588 |
| PDGF_BB_14 | intron      | ACTR1A        | 10 | 104260388 |
| PDGF_BB_15 | 3_prime_UTR | TMEM120B      | 12 | 122216910 |
| PDGF_BB_15 | intron      | HPD           | 12 | 122289308 |
| PDGF_BB_16 | upstream    | PEAR1         | 1  | 156862468 |
| PDGF_BB_16 | intron      | PEAR1         | 1  | 156867990 |
| PDGF_BB_16 | intron      | PEAR1         | 1  | 156869630 |
| PDGF_BB_16 | synonymous  | PEAR1         | 1  | 156879580 |
| PDGF_BB_17 | NA          | NA            | 14 | 105727251 |
| PDGF_BB_17 | intron      | BRF1          | 14 | 105737261 |
| PDGF_BB_19 | intron      | TSPOAP1       | 17 | 56398479  |
| PDGF_BB_19 | NA          | NA            | NA | NA        |
| PDGF_BB_2  | NA          | NA            | 6  | 163753830 |
| PDGF_BB_2  | NA          | NA            | 6  | 163838794 |
| PDGF_BB_2  | NA          | NA            | 6  | 163864331 |
| PDGF_BB_3  | intron      | SLC24A3       | 20 | 19285511  |
| PDGF_BB_3  | intron      | SLC24A3       | 20 | 19321537  |
| PDGF_BB_4  | intron      | SLC8A3        | 14 | 70623306  |
| PDGF_BB_4  | intron      | SLC8A3        | 14 | 70653758  |
| PDGF_BB_5  | intron      | NFASC         | 1  | 204956325 |
| PDGF_BB_5  | intron      | NFASC         | 1  | 204982054 |
| PDGF_BB_5  | intergenic  | NFASC         | 1  | 205000315 |
| PDGF_BB_5  | -           | -             | 1  | 205017962 |

|           |             |               |    |           |
|-----------|-------------|---------------|----|-----------|
| PDGF_BB_5 | intron      | CNTN2         | 1  | 205040943 |
| PDGF_BB_5 | downstream  | CNTN2         | 1  | 205049383 |
| PDGF_BB_5 | intron      | RBBP5         | 1  | 205065298 |
| PDGF_BB_5 | intron      | RBBP5         | 1  | 205070178 |
| PDGF_BB_5 | intron      | RBBP5         | 1  | 205078247 |
| PDGF_BB_5 | intron      | RBBP5         | 1  | 205081380 |
| PDGF_BB_5 | intron      | RBBP5         | 1  | 205084120 |
| PDGF_BB_5 | intron      | RBBP5         | 1  | 205090157 |
| PDGF_BB_5 | upstream    | RP11-383G10.1 | 1  | 205099959 |
| PDGF_BB_5 | downstream  | RP11-383G10.1 | 1  | 205104550 |
| PDGF_BB_5 | downstream  | DSTYK         | 1  | 205108083 |
| PDGF_BB_5 | intron      | DSTYK         | 1  | 205120640 |
| PDGF_BB_5 | intron      | DSTYK         | 1  | 205123021 |
| PDGF_BB_7 | intron      | SERPINE2      | 2  | 224874874 |
| PDGF_BB_7 | intron      | SERPINE2      | 2  | 224880498 |
| PDGF_BB_7 | intergenic  | SERPINE2      | 2  | 224923426 |
| PDGF_BB_7 | intergenic  | SERPINE2      | 2  | 224942369 |
| PDGF_BB_9 | intron      | PCSK6         | 15 | 101985142 |
| PDGF_BB_9 | intron      | PCSK6         | 15 | 101992998 |
| RANTES_2  | intergenic  | ANKRD30A      | 10 | 37693541  |
| RANTES_2  | intergenic  | TACC1P1       | 10 | 37766696  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56820356  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56849055  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56849749  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56858890  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56920033  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56923041  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56925884  |
| RANTES_4  | intron      | ARHGEF3       | 3  | 56942411  |
| RANTES_8  | intron      | HEATR9        | 17 | 34189017  |
| RANTES_8  | 5_prime_UTR | CCL5          | 17 | 34207780  |
| SCF_1     | intron      | RANBP10       | 16 | 67803063  |
| SCF_1     | intron      | RANBP10       | 16 | 67835177  |
| SCF_1     | intron      | TSNAXIP1      | 16 | 67861134  |
| SCF_10    | NA          | NA            | 20 | 44503605  |
| SCF_10    | intron      | PLTP          | 20 | 44534651  |
| SCF_10    | intron      | PLTP          | 20 | 44536932  |
| SCF_10    | upstream    | PLTP          | 20 | 44545460  |
| SCF_10    | upstream    | PLTP          | 20 | 44545773  |
| SCF_10    | intergenic  | RP11-323C15.1 | 20 | 44548301  |
| SCF_10    | NA          | NA            | 20 | 44551600  |
| SCF_10    | intergenic  | RP11-323C15.1 | 20 | 44557474  |
| SCF_10    | NA          | NA            | 20 | 44563007  |
| SCF_10    | NA          | NA            | 20 | 44596545  |
| SCF_10    | NA          | NA            | 20 | 44615889  |
| SCF_10    | intergenic  | RP11-323C15.1 | 20 | 44624117  |

|        |            |               |    |           |
|--------|------------|---------------|----|-----------|
| SCF_10 | intergenic | RP11-323C15.1 | 20 | 44628141  |
| SCF_10 | NA         | NA            | NA | NA        |
| SCF_12 | intron     | TTC39B        | 9  | 15201410  |
| SCF_12 | missense   | TTC39B        | 9  | 15214237  |
| SCF_12 | intron     | TTC39B        | 9  | 15227761  |
| SCF_12 | intron     | TTC39B        | 9  | 15272064  |
| SCF_12 | intron     | TTC39B        | 9  | 15273257  |
| SCF_12 | intron     | TTC39B        | 9  | 15275582  |
| SCF_12 | intron     | TTC39B        | 9  | 15285448  |
| SCF_12 | intron     | TTC39B        | 9  | 15288198  |
| SCF_12 | intron     | TTC39B        | 9  | 15289674  |
| SCF_12 | intron     | TTC39B        | 9  | 15293427  |
| SCF_12 | intron     | TTC39B        | 9  | 15296034  |
| SCF_12 | intron     | TTC39B        | 9  | 15296845  |
| SCF_12 | intron     | TTC39B        | 9  | 15297629  |
| SCF_12 | intron     | TTC39B        | 9  | 15298666  |
| SCF_12 | intron     | TTC39B        | 9  | 15300849  |
| SCF_12 | intron     | TTC39B        | 9  | 15301428  |
| SCF_12 | intron     | TTC39B        | 9  | 15302613  |
| SCF_12 | -          | TTC39B        | 9  | 15303632  |
| SCF_12 | intron     | TTC39B        | 9  | 15304782  |
| SCF_12 | intron     | TTC39B        | 9  | 15306292  |
| SCF_12 | upstream   | TTC39B        | 9  | 15307746  |
| SCF_12 | intergenic | TTC39B        | 9  | 15327379  |
| SCF_12 | intergenic | RN7SL157P     | 9  | 15328258  |
| SCF_12 | intergenic | RN7SL157P     | 9  | 15330506  |
| SCF_12 | intergenic | RN7SL157P     | 9  | 15340597  |
| SCF_12 | upstream   | RN7SL157P     | 9  | 15343391  |
| SCF_12 | intergenic | SNAPC3        | 9  | 15393875  |
| SCF_12 | intergenic | SNAPC3        | 9  | 15397244  |
| SCF_12 | intergenic | SNAPC3        | 9  | 15397743  |
| SCF_13 | intron     | ABCA1         | 9  | 107585213 |
| SCF_13 | intron     | ABCA1         | 9  | 107657070 |
| SCF_14 | intron     | AXDND1        | 1  | 179441653 |
| SCF_14 | missense   | NPHS2         | 1  | 179526214 |
| SCF_14 | synonymous | NPHS2         | 1  | 179544898 |
| SCF_14 | intron     | TDRD5         | 1  | 179571862 |
| SCF_17 | intron     | PDILT         | 16 | 20377587  |
| SCF_17 | intron     | PDILT         | 16 | 20400839  |
| SCF_18 | intron     | AFF1          | 4  | 88009779  |
| SCF_18 | intron     | AFF1          | 4  | 88030261  |
| SCF_19 | intergenic | TTBK2         | 15 | 43219196  |
| SCF_19 | intron     | UBR1          | 15 | 43244141  |
| SCF_19 | intron     | UBR1          | 15 | 43278726  |
| SCF_19 | intron     | UBR1          | 15 | 43285629  |
| SCF_19 | intron     | UBR1          | 15 | 43297021  |

|        |             |               |    |           |
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| SCF_19 | intron      | UBR1          | 15 | 43297183  |
| SCF_19 | intron      | UBR1          | 15 | 43341747  |
| SCF_2  | upstream    | CETP          | 16 | 56993161  |
| SCF_2  | -           | -             | 16 | 56996158  |
| SCF_20 | intron      | TOMM40L       | 1  | 161197244 |
| SCF_20 | intron      | NR1I3         | 1  | 161204426 |
| SCF_21 | intron      | PPP1R9A       | 7  | 94909495  |
| SCF_21 | 3_prime_UTR | PPP1R9A       | 7  | 94925614  |
| SCF_21 | intron      | PON1          | 7  | 94945215  |
| SCF_21 | intron      | PON1          | 7  | 94945872  |
| SCF_21 | NA          | NA            | 7  | 94946084  |
| SCF_21 | intron      | PON1          | 7  | 94952816  |
| SCF_21 | intergenic  | PON1          | 7  | 94961567  |
| SCF_21 | downstream  | PON3          | 7  | 94985225  |
| SCF_21 | intron      | PON3          | 7  | 94995519  |
| SCF_21 | intron      | PON3          | 7  | 95015586  |
| SCF_22 | intergenic  | NSMCE2        | 8  | 126384737 |
| SCF_22 | intergenic  | NSMCE2        | 8  | 126385396 |
| SCF_22 | intergenic  | NSMCE2        | 8  | 126409645 |
| SCF_22 | intron      | TRIB1         | 8  | 126447473 |
| SCF_22 | intergenic  | RN7SL590P     | 8  | 126465832 |
| SCF_22 | upstream    | RP11-136O12.2 | 8  | 126476873 |
| SCF_22 | intron      | RP11-136O12.2 | 8  | 126479362 |
| SCF_22 | intron      | RP11-136O12.2 | 8  | 126480972 |
| SCF_22 | intron      | RP11-136O12.2 | 8  | 126486409 |
| SCF_22 | intron      | RP11-136O12.2 | 8  | 126550180 |
| SCF_23 | intron      | GALNT2        | 1  | 230297136 |
| SCF_23 | intron      | GALNT2        | 1  | 230297778 |
| SCF_6  | intron      | ALDH1A2       | 15 | 58581426  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58582195  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58587409  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58605881  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58611396  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58611859  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58612816  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58633380  |
| SCF_6  | -           | -             | 15 | 58635638  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58639710  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58644087  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58651580  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58652220  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58659418  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58659761  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58670135  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58674695  |
| SCF_6  | intron      | ALDH1A2       | 15 | 58676821  |

|       |            |             |    |           |
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| SCF_6 | intron     | ALDH1A2     | 15 | 58680638  |
| SCF_6 | -          | -           | 15 | 58680639  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58680643  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58680954  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58681509  |
| SCF_6 | -          | -           | 15 | 58683366  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58683542  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58685970  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58686157  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58686540  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58687932  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58695948  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58696300  |
| SCF_6 | intron     | ALDH1A2     | 15 | 58698421  |
| SCF_6 | intron     | LIPC        | 15 | 58706926  |
| SCF_6 | intron     | LIPC        | 15 | 58716044  |
| SCF_6 | intron     | LIPC        | 15 | 58718998  |
| SCF_6 | intron     | LIPC        | 15 | 58719384  |
| SCF_6 | intron     | LIPC        | 15 | 58723939  |
| SCF_6 | intron     | LIPC        | 15 | 58724706  |
| SCF_6 | intron     | LIPC        | 15 | 58727325  |
| SCF_6 | intron     | LIPC        | 15 | 58728878  |
| SCF_6 | intron     | LIPC        | 15 | 58730498  |
| SCF_6 | intron     | LIPC        | 15 | 58731818  |
| SCF_6 | intron     | LIPC        | 15 | 58739761  |
| SCF_6 | intron     | LIPC        | 15 | 58741070  |
| SCF_6 | intron     | LIPC        | 15 | 58741384  |
| SCF_6 | intron     | LIPC        | 15 | 58741891  |
| SCF_6 | intron     | LIPC        | 15 | 58750118  |
| SCF_6 | intron     | LIPC        | 15 | 58762377  |
| SCF_6 | intron     | LIPC        | 15 | 58772815  |
| SCF_6 | NA         | NA          | NA | NA        |
| SCF_7 | NA         | NA          | 1  | 234826267 |
| SCF_7 | NA         | NA          | 1  | 234858597 |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89285292  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89287024  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89292082  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89294185  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89302702  |
| SCF_9 | NA         | NA          | 12 | 89338612  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89340293  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89368688  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89369501  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89377792  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89381246  |
| SCF_9 | intergenic | RP11-13A1.1 | 12 | 89383136  |

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| SCF_9     | -                          | -             | 12 | 89383147  |
| SCF_9     | intergenic                 | RP11-13A1.1   | 12 | 89383706  |
| SCF_9     | intergenic                 | RP11-13A1.1   | 12 | 89384112  |
| SCF_9     | intergenic                 | RP11-13A1.1   | 12 | 89385017  |
| SCF_9     | intergenic                 | RP11-13A1.1   | 12 | 89387163  |
| SCF_9     | intergenic                 | RP11-13A1.1   | 12 | 89387344  |
| SCF_9     | intergenic                 | RP11-13A1.1   | 12 | 89388031  |
| SCF_9     | non_coding_transcript_exon | RP11-13A1.1   | 12 | 89408584  |
| SCF_9     | upstream                   | RP11-13A1.1   | 12 | 89413835  |
| SCF_9     | upstream                   | RP11-13A1.1   | 12 | 89417009  |
| SCF_9     | intron                     | RP11-13A1.3   | 12 | 89450901  |
| SCGF_B_1  | intergenic                 | MRPL3         | 3  | 128306323 |
| SCGF_B_1  | intron                     | RPN1          | 3  | 128383539 |
| SCGF_B_13 | upstream                   | GCNT1         | 9  | 79033198  |
| SCGF_B_13 | intron                     | GCNT1         | 9  | 79110078  |
| SCGF_B_13 | missense                   | GCNT1         | 9  | 79117751  |
| SCGF_B_13 | 3_prime_UTR                | GCNT1         | 9  | 79120740  |
| SCGF_B_14 | upstream                   | RPSAP5        | 14 | 103843344 |
| SCGF_B_14 | intron                     | MARK3         | 14 | 103904842 |
| SCGF_B_16 | intron                     | SLC5A8        | 12 | 101598983 |
| SCGF_B_16 | intergenic                 | SLC5A8        | 12 | 101611442 |
| SCGF_B_17 | intron                     | WDFY2         | 13 | 52178166  |
| SCGF_B_17 | intron                     | WDFY2         | 13 | 52223081  |
| SCGF_B_18 | intron                     | STAB2         | 12 | 104058214 |
| SCGF_B_18 | intron                     | STAB2         | 12 | 104084581 |
| SCGF_B_18 | intron                     | STAB2         | 12 | 104144835 |
| SCGF_B_18 | intron                     | STAB2         | 12 | 104149022 |
| SCGF_B_18 | intron                     | STAB2         | 12 | 104149874 |
| SCGF_B_18 | intron                     | TTC41P        | 12 | 104241404 |
| SCGF_B_19 | intron                     | SLCO5A1       | 8  | 70727663  |
| SCGF_B_19 | intron                     | SLCO5A1       | 8  | 70732155  |
| SCGF_B_22 | intron                     | SYT3          | 19 | 51137442  |
| SCGF_B_22 | intron                     | SYT3          | 19 | 51140506  |
| SCGF_B_22 | upstream                   | C19orf81      | 19 | 51150548  |
| SCGF_B_22 | intron                     | SHANK1        | 19 | 51181589  |
| SCGF_B_22 | intron                     | SHANK1        | 19 | 51222296  |
| SCGF_B_22 | upstream                   | SHANK1        | 19 | 51223567  |
| SCGF_B_22 | downstream                 | CLEC11A       | 19 | 51232864  |
| SCGF_B_22 | downstream                 | CLEC11A       | 19 | 51233110  |
| SCGF_B_22 | intergenic                 | CLEC11A       | 19 | 51234285  |
| SCGF_B_22 | intergenic                 | CLEC11A       | 19 | 51239179  |
| SCGF_B_22 | intron                     | CTD-2568A17.1 | 19 | 51291533  |
| SCGF_B_26 | intron                     | PCSK2         | 20 | 17460471  |
| SCGF_B_26 | -                          | -             | 20 | 17471551  |
| SCGF_B_26 | intron                     | BFSP1         | 20 | 17523651  |
| SCGF_B_26 | intron                     | DSTN          | 20 | 17552349  |



|           |             |               |    |           |
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| SCGF_B_26 | intron      | DSTN          | 20 | 17555980  |
| SCGF_B_26 | intron      | DSTN          | 20 | 17556020  |
| SCGF_B_26 | intron      | DSTN          | 20 | 17567578  |
| SCGF_B_26 | intron      | RRBP1         | 20 | 17595329  |
| SCGF_B_26 | missense    | RRBP1         | 20 | 17596155  |
| SCGF_B_26 | intron      | RRBP1         | 20 | 17596731  |
| SCGF_B_26 | intron      | RRBP1         | 20 | 17599752  |
| SCGF_B_26 | intron      | RRBP1         | 20 | 17602028  |
| SCGF_B_26 | -           | RRBP1         | 20 | 17605016  |
| SCGF_B_26 | intron      | RRBP1         | 20 | 17606372  |
| SCGF_B_27 | intron      | MFSD6         | 2  | 191286658 |
| SCGF_B_27 | intron      | MFSD6         | 2  | 191313902 |
| SCGF_B_28 | intergenic  | RP11-80A15.1  | 14 | 24951160  |
| SCGF_B_28 | intron      | CMA1          | 14 | 24975032  |
| SCGF_B_28 | intergenic  | RP11-104E19.1 | 14 | 25056924  |
| SCGF_B_28 | upstream    | RP11-104E19.1 | 14 | 25064826  |
| SCGF_B_28 | downstream  | GZMH          | 14 | 25070800  |
| SCGF_B_28 | downstream  | GZMH          | 14 | 25075531  |
| SCGF_B_28 | upstream    | GZMH          | 14 | 25080289  |
| SCGF_B_28 | intron      | GZMB          | 14 | 25101397  |
| SCGF_B_28 | intron      | RP11-104E19.1 | 14 | 25118736  |
| SCGF_B_28 | downstream  | RP11-104E19.1 | 14 | 25128403  |
| SCGF_B_29 | intron      | CCDC26        | 8  | 130569893 |
| SCGF_B_29 | intron      | CCDC26        | 8  | 130589847 |
| SCGF_B_29 | intron      | CCDC26        | 8  | 130589974 |
| SCGF_B_29 | intron      | CCDC26        | 8  | 130603635 |
| SCGF_B_3  | intron      | ATP8B4        | 15 | 50357743  |
| SCGF_B_3  | intron      | ATP8B4        | 15 | 50374529  |
| SCGF_B_30 | intron      | CTD-2631K10.1 | 5  | 72090894  |
| SCGF_B_30 | intron      | TNPO1         | 5  | 72139788  |
| SCGF_B_30 | intron      | TNPO1         | 5  | 72154551  |
| SCGF_B_30 | intron      | TNPO1         | 5  | 72180647  |
| SCGF_B_30 | intron      | TNPO1         | 5  | 72197104  |
| SCGF_B_30 | upstream    | FCHO2         | 5  | 72248719  |
| SCGF_B_32 | intron      | MYBPC2        | 19 | 50955876  |
| SCGF_B_32 | missense    | MYBPC2        | 19 | 50957600  |
| SCGF_B_32 | intron      | MYBPC2        | 19 | 50965990  |
| SCGF_B_32 | intron      | MYBPC2        | 19 | 50968890  |
| SCGF_B_32 | intron      | FAM71E1       | 19 | 50972522  |
| SCGF_B_32 | intron      | FAM71E1       | 19 | 50973259  |
| SCGF_B_32 | intron      | FAM71E1       | 19 | 50976882  |
| SCGF_B_32 | intron      | FAM71E1       | 19 | 50977689  |
| SCGF_B_32 | intron      | EMC10         | 19 | 50982086  |
| SCGF_B_32 | 3_prime_UTR | EMC10         | 19 | 50987992  |
| SCGF_B_32 | 3_prime_UTR | EMC10         | 19 | 50988003  |
| SCGF_B_32 | 3_prime_UTR | EMC10         | 19 | 50989217  |

|           |               |             |    |           |
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| SCGF_B_32 | downstream    | EMC10       | 19 | 50995880  |
| SCGF_B_32 | downstream    | EMC10       | 19 | 50997246  |
| SCGF_B_32 | downstream    | JOSD2       | 19 | 51008877  |
| SCGF_B_32 | intron        | JOSD2       | 19 | 51010653  |
| SCGF_B_32 | intron        | JOSD2       | 19 | 51012827  |
| SCGF_B_32 | upstream      | ASPDH       | 19 | 51018813  |
| SCGF_B_32 | intron        | LRRC4B      | 19 | 51025819  |
| SCGF_B_32 | intron        | LRRC4B      | 19 | 51029238  |
| SCGF_B_32 | intron        | LRRC4B      | 19 | 51030134  |
| SCGF_B_32 | intron        | LRRC4B      | 19 | 51038708  |
| SCGF_B_32 | intron        | LRRC4B      | 19 | 51041781  |
| SCGF_B_32 | intron        | LRRC4B      | 19 | 51045095  |
| SCGF_B_32 | intron        | LRRC4B      | 19 | 51053661  |
| SCGF_B_32 | upstream      | LRRC4B      | 19 | 51073639  |
| SCGF_B_32 | intergenic    | LRRC4B      | 19 | 51077038  |
| SCGF_B_32 | intergenic    | LRRC4B      | 19 | 51089778  |
| SCGF_B_32 | intergenic    | LRRC4B      | 19 | 51091010  |
| SCGF_B_32 | intergenic    | LRRC4B      | 19 | 51091617  |
| SCGF_B_32 | intergenic    | SYT3        | 19 | 51098368  |
| SCGF_B_32 | intergenic    | SYT3        | 19 | 51110830  |
| SCGF_B_32 | intergenic    | SYT3        | 19 | 51113526  |
| SCGF_B_32 | synonymous    | SYT3        | 19 | 51128661  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51129745  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51129919  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51130144  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51130554  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51130909  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51134973  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51135051  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51135457  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51136844  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51137292  |
| SCGF_B_32 | splice_region | SYT3        | 19 | 51140689  |
| SCGF_B_32 | 5_prime_UTR   | SYT3        | 19 | 51140752  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51141334  |
| SCGF_B_32 | intron        | SYT3        | 19 | 51142845  |
| SCGF_B_32 | upstream      | C19orf81    | 19 | 51150249  |
| SCGF_B_7  | intron        | PROSER2     | 10 | 11895507  |
| SCGF_B_7  | intron        | PROSER2-AS1 | 10 | 11929674  |
| SCGF_B_8  | intron        | RAB11FIP3   | 16 | 531056    |
| SCGF_B_8  | intron        | RAB11FIP3   | 16 | 540468    |
| SCGF_B_8  | intron        | RAB11FIP3   | 16 | 541671    |
| SCGF_B_8  | intron        | RAB40C      | 16 | 643064    |
| SCGF_B_9  | intergenic    | RNU6-968P   | 1  | 236086104 |
| SCGF_B_9  | intergenic    | RNU6-968P   | 1  | 236087992 |
| SDF_1A_1  | intergenic    | LEPR        | 1  | 66135991  |

|           |            |           |    |           |
|-----------|------------|-----------|----|-----------|
| SDF_1A_1  | intergenic | LEPR      | 1  | 66159278  |
| SDF_1A_11 | intron     | PPARG     | 3  | 12442731  |
| SDF_1A_11 | intergenic | LINC00690 | 3  | 12517103  |
| SDF_1A_11 | intergenic | LINC00690 | 3  | 12519202  |
| SDF_1A_11 | intron     | TSEN2     | 3  | 12538595  |
| SDF_1A_11 | intron     | TSEN2     | 3  | 12547147  |
| SDF_1A_11 | upstream   | MKRN2     | 3  | 12596333  |
| SDF_1A_12 | intergenic | DPP6      | 7  | 153557997 |
| SDF_1A_12 | NA         | NA        | 7  | 153568661 |
| SDF_1A_12 | intergenic | DPP6      | 7  | 153571324 |
| SDF_1A_12 | upstream   | DPP6      | 7  | 153581933 |
| SDF_1A_12 | NA         | NA        | 7  | 153587207 |
| SDF_1A_12 | NA         | NA        | 7  | 153591994 |
| SDF_1A_12 | NA         | NA        | 7  | 153593813 |
| SDF_1A_12 | NA         | NA        | 7  | 153595436 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153598592 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153605934 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153606635 |
| SDF_1A_12 | NA         | NA        | 7  | 153609061 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153617821 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153618948 |
| SDF_1A_12 | NA         | NA        | 7  | 153619062 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153621035 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153624973 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153632459 |
| SDF_1A_12 | NA         | NA        | 7  | 153637111 |
| SDF_1A_12 | -          | -         | 7  | 153637982 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153638317 |
| SDF_1A_12 | NA         | NA        | 7  | 153649423 |
| SDF_1A_12 | NA         | NA        | 7  | 153650668 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153655480 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153657480 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153658104 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153660141 |
| SDF_1A_12 | NA         | NA        | 7  | 153660427 |
| SDF_1A_12 | NA         | NA        | 7  | 153670448 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153677011 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153699295 |
| SDF_1A_12 | intron     | DPP6      | 7  | 153707385 |
| SDF_1A_12 | NA         | NA        | 7  | 153710624 |
| SDF_1A_12 | NA         | NA        | 7  | 153721613 |
| SDF_1A_12 | NA         | NA        | 7  | 153723760 |
| SDF_1A_12 | NA         | NA        | NA | NA        |
| SDF_1A_13 | NA         | NA        | 5  | 115298232 |
| SDF_1A_13 | NA         | NA        | 5  | 115351224 |
| SDF_1A_14 | NA         | NA        | 17 | 66989106  |

|           |             |            |    |           |
|-----------|-------------|------------|----|-----------|
| SDF_1A_14 | intron      | ABCA9      | 17 | 67003528  |
| SDF_1A_2  | intergenic  | CXCL12     | 10 | 44840850  |
| SDF_1A_2  | intergenic  | CXCL12     | 10 | 44857244  |
| SDF_1A_2  | upstream    | CXCL12     | 10 | 44881660  |
| SDF_1A_2  | upstream    | CXCL12     | 10 | 44882201  |
| SDF_1A_2  | intergenic  | CXCL12     | 10 | 44893756  |
| SDF_1A_3  | intergenic  | CXCL12     | 10 | 44829089  |
| SDF_1A_3  | intron      | CXCL12     | 10 | 44878114  |
| SDF_1A_3  | upstream    | CXCL12     | 10 | 44886605  |
| SDF_1A_3  | intergenic  | CXCL12     | 10 | 44893756  |
| SDF_1A_4  | intron      | SLC4A10    | 2  | 162822349 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162852079 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162859000 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162866992 |
| SDF_1A_4  | NA          | NA         | 2  | 162872677 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162877028 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162887969 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162894025 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162895963 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162901327 |
| SDF_1A_4  | NA          | NA         | 2  | 162908664 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162910403 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162910536 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162913498 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162916578 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162917337 |
| SDF_1A_4  | NA          | NA         | 2  | 162920003 |
| SDF_1A_4  | NA          | NA         | 2  | 162921606 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162922728 |
| SDF_1A_4  | intron      | DPP4       | 2  | 162923917 |
| SDF_1A_4  | NA          | NA         | 2  | 162925618 |
| SDF_1A_4  | 5_prime_UTR | DPP4       | 2  | 162930725 |
| SDF_1A_4  | upstream    | DPP4       | 2  | 162933561 |
| SDF_1A_4  | upstream    | TIMM8AP1   | 2  | 162938654 |
| SDF_1A_4  | upstream    | TIMM8AP1   | 2  | 162938911 |
| SDF_1A_4  | intergenic  | EIF3EP2    | 2  | 162939467 |
| SDF_1A_4  | NA          | NA         | 2  | 162952349 |
| SDF_1A_4  | intergenic  | AC008063.3 | 2  | 162958136 |
| SDF_1A_4  | intergenic  | AC008063.3 | 2  | 162960210 |
| SDF_1A_4  | intergenic  | AC008063.3 | 2  | 162964301 |
| SDF_1A_4  | intergenic  | GCG        | 2  | 162977965 |
| SDF_1A_4  | intergenic  | GCG        | 2  | 162982138 |
| SDF_1A_4  | intergenic  | GCG        | 2  | 162990608 |
| SDF_1A_4  | intergenic  | GCG        | 2  | 162991075 |
| SDF_1A_4  | downstream  | GCG        | 2  | 162996174 |
| SDF_1A_4  | NA          | NA         | 2  | 162999388 |

|          |            |               |    |           |
|----------|------------|---------------|----|-----------|
| SDF_1A_4 | NA         | NA            | 2  | 163007575 |
| SDF_1A_4 | upstream   | GCG           | 2  | 163012730 |
| SDF_1A_6 | downstream | RP11-395D3.1  | 9  | 89315229  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89322570  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89332996  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89340501  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89343747  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89351788  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89356766  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89368334  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89371107  |
| SDF_1A_6 | intergenic | RP11-395D3.1  | 9  | 89435024  |
| TNF_B_2  | NA         | NA            | 1  | 196657414 |
| TNF_B_2  | NA         | NA            | 1  | 196669581 |
| TNF_B_2  | NA         | NA            | 1  | 196672385 |
| TNF_B_2  | NA         | NA            | NA | NA        |
| TRAIL_10 | downstream | AHSG          | 3  | 186339617 |
| TRAIL_10 | intron     | RP11-573D15.8 | 3  | 186420948 |
| TRAIL_12 | intron     | TRAPPC8       | 18 | 29502140  |
| TRAIL_12 | intron     | TRAPPC8       | 18 | 29515691  |
| TRAIL_12 | upstream   | TRAPPC8       | 18 | 29535266  |
| TRAIL_12 | downstream | RP11-326K13.3 | 18 | 29569351  |
| TRAIL_12 | downstream | RP11-326K13.3 | 18 | 29569416  |
| TRAIL_12 | intergenic | RNF125        | 18 | 29586371  |
| TRAIL_12 | downstream | RNF125        | 18 | 29655791  |
| TRAIL_12 | intron     | RP11-53I6.2   | 18 | 29660305  |
| TRAIL_12 | intron     | RP11-53I6.2   | 18 | 29666127  |
| TRAIL_5  | intron     | TNFSF10       | 3  | 172234017 |
| TRAIL_5  | upstream   | TNFSF10       | 3  | 172242764 |
| TRAIL_5  | intergenic | U8            | 3  | 172264728 |
| TRAIL_5  | intergenic | U8            | 3  | 172265633 |
| TRAIL_7  | intron     | FNDC3B        | 3  | 172075962 |
| TRAIL_7  | intron     | FNDC3B        | 3  | 172078051 |
| TRAIL_7  | intron     | FNDC3B        | 3  | 172091227 |
| TRAIL_7  | intron     | FNDC3B        | 3  | 172111829 |
| TRAIL_7  | -          | -             | 3  | 172118997 |
| TRAIL_7  | downstream | FNDC3B        | 3  | 172119811 |
| TRAIL_7  | intergenic | U8            | 3  | 172129815 |
| TRAIL_7  | intergenic | U8            | 3  | 172156117 |
| TRAIL_7  | downstream | GHSR          | 3  | 172158924 |
| TRAIL_7  | downstream | GHSR          | 3  | 172162829 |
| TRAIL_7  | synonymous | GHSR          | 3  | 172165757 |
| TRAIL_7  | intergenic | U8            | 3  | 172200214 |
| TRAIL_7  | intergenic | U8            | 3  | 172211182 |
| TRAIL_7  | intergenic | U8            | 3  | 172211743 |
| TRAIL_7  | intergenic | U8            | 3  | 172218253 |

|         |            |              |    |           |
|---------|------------|--------------|----|-----------|
| TRAIL_7 | downstream | TNFSF10      | 3  | 172221003 |
| TRAIL_7 | intron     | TNFSF10      | 3  | 172234017 |
| TRAIL_7 | intron     | TNFSF10      | 3  | 172238420 |
| TRAIL_7 | intron     | TNFSF10      | 3  | 172238498 |
| TRAIL_7 | upstream   | TNFSF10      | 3  | 172241866 |
| TRAIL_7 | upstream   | TNFSF10      | 3  | 172243161 |
| TRAIL_7 | intergenic | U8           | 3  | 172246664 |
| TRAIL_7 | intergenic | U8           | 3  | 172252149 |
| TRAIL_7 | intergenic | U8           | 3  | 172256054 |
| TRAIL_7 | intergenic | U8           | 3  | 172256154 |
| TRAIL_7 | intergenic | U8           | 3  | 172256714 |
| TRAIL_9 | missense   | SERPINA1     | 14 | 94844947  |
| TRAIL_9 | intron     | SERPINA1     | 14 | 94846892  |
| VEGF_12 | intron     | VLDLR-AS1    | 9  | 2536666   |
| VEGF_12 | intron     | VLDLR-AS1    | 9  | 2543316   |
| VEGF_12 | intron     | VLDLR        | 9  | 2638113   |
| VEGF_12 | intergenic | VLDLR        | 9  | 2674729   |
| VEGF_12 | intergenic | VLDLR        | 9  | 2680425   |
| VEGF_12 | intergenic | VLDLR        | 9  | 2682943   |
| VEGF_12 | intergenic | VLDLR        | 9  | 2686273   |
| VEGF_12 | intergenic | VLDLR        | 9  | 2686336   |
| VEGF_12 | intergenic | KCNV2        | 9  | 2692583   |
| VEGF_12 | intergenic | KCNV2        | 9  | 2694290   |
| VEGF_12 | intergenic | KCNV2        | 9  | 2694834   |
| VEGF_3  | downstream | VEGFA        | 6  | 43756861  |
| VEGF_3  | downstream | VEGFA        | 6  | 43757896  |
| VEGF_3  | upstream   | RP11-344J7.3 | 6  | 43766929  |
| VEGF_3  | intron     | RP11-344J7.3 | 6  | 43788415  |
| VEGF_3  | intron     | RP11-344J7.3 | 6  | 43790318  |
| VEGF_3  | intron     | RP11-344J7.4 | 6  | 43815148  |
| VEGF_3  | intron     | RP11-344J7.4 | 6  | 43819046  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43828477  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43832966  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43845044  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43849964  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43855489  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43887461  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43892269  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43900707  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43902928  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43911598  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43913243  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43913988  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43916310  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43919416  |
| VEGF_3  | intergenic | ACTG1P9      | 6  | 43919650  |

|        |            |         |    |           |
|--------|------------|---------|----|-----------|
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43919695  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43922603  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43922826  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43924763  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43924774  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43924830  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43925526  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43925778  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43926219  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43926635  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43926638  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43928108  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43928261  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43929421  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43930985  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43932319  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43932612  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43932763  |
| VEGF_3 | -          | -       | 6  | 43939313  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43946490  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43949907  |
| VEGF_3 | intergenic | ACTG1P9 | 6  | 43952089  |
| VEGF_6 | intron     | MEF2C   | 5  | 88098064  |
| VEGF_6 | intron     | MEF2C   | 5  | 88128642  |
| VEGF_7 | intron     | ZFPM1   | 16 | 88571196  |
| VEGF_7 | intron     | ZFPM1   | 16 | 88573453  |
| VEGF_9 | intron     | ZFPM2   | 8  | 106471906 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106517072 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106519305 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106521845 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106524285 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106527975 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106528030 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106531668 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106531674 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106532477 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106534599 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106536267 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106559719 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106570222 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106574238 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106583124 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106583528 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106592539 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106593207 |
| VEGF_9 | intron     | ZFPM2   | 8  | 106604227 |





[illegible]

[illegible]

|               |             |
|---------------|-------------|
| intergenic    | SLC2A2      |
| intergenic    | SLC2A2      |
| intron        | PSG8        |
| intron        | PSG6        |
| intron        | IRGC        |
| intron        | KCNN4       |
| intron        | KCNN4       |
| -             | -           |
| downstream    | LYPD5       |
| intron        | LYPD5       |
| intron        | LYPD5       |
| intron        | LYPD5       |
| intron        | LYPD5       |
| intron        | LYPD5       |
| intron        | LYPD5       |
| intron        | LYPD5       |
| splice_region | ZNF283      |
| 3_prime_UTR   | ZNF283      |
| intron        | ZNF283      |
| intron        | ZNF283      |
| intron        | ZNF404      |
| intron        | RP11-15A1.2 |
| intron        | ZNF229      |
| upstream      | ZNF229      |
| downstream    | ZNF180      |
| synonymous    | ZNF180      |
| intron        | ZNF180      |
| NA            | NA          |
| intron        | ZNF180      |
| intron        | ZNF180      |
| NA            | NA          |
| intron        | CEACAM20    |
| intron        | CEACAM20    |
| synonymous    | CEACAM20    |
| intron        | CEACAM20    |
| upstream      | CEACAM20    |
| intron        | CEACAM22P   |
| intron        | CEACAM22P   |
| intron        | CEACAM22P   |
| intron        | CEACAM22P   |
| intron        | CEACAM22P   |
| intron        | CEACAM22P   |
| intron        | CEACAM22P   |
| NA            | NA          |
| intron        | CEACAM22P   |
| 5_prime_UTR   | IGSF23      |

|             |               |
|-------------|---------------|
| intron      | IGSF23        |
| intron      | IGSF23        |
| NA          | NA            |
| intron      | IGSF23        |
| intergenic  | snoZ6         |
| intron      | CBLC          |
| intron      | BCAM          |
| downstream  | BCAM          |
| intergenic  | BCAM          |
| intergenic  | NECTIN2       |
| upstream    | NECTIN2       |
| upstream    | NECTIN2       |
| intron      | NECTIN2       |
| intron      | NECTIN2       |
| intron      | TOMM40        |
| intron      | TOMM40        |
| intron      | APOC1         |
| downstream  | APOC1         |
| intron      | OPA3          |
| 3_prime_UTR | OPA3          |
| intron      | EML2          |
| intron      | EML2          |
| -           | -             |
| -           | -             |
| downstream  | RN7SL836P     |
| downstream  | RN7SL836P     |
| intergenic  | RN7SL836P     |
| upstream    | GIPR          |
| intron      | GIPR          |
| intron      | GIPR          |
| intron      | QPCTL         |
| intron      | FBXO46        |
| NA          | NA            |
| NA          | NA            |
| NA          | NA            |
| intergenic  | AC112518.3    |
| intergenic  | AC112518.3    |
| intron      | RP11-642E20.3 |
| downstream  | YTHDC1        |
| downstream  | YTHDC1        |
| intron      | YTHDC1        |
| intron      | YTHDC1        |
| intron      | YTHDC1        |
| intron      | YTHDC1        |
| intergenic  | AC112518.3    |
| intergenic  | AC112518.3    |

|            |            |
|------------|------------|
| intergenic | AC112518.3 |
| downstream | MT2P1      |
| downstream | MT2P1      |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intron     | AMBN       |
| intron     | RUFY3      |
| intron     | TTC21A     |
| intron     | TTC21A     |
| intron     | NPFFR2     |
| intron     | NPFFR2     |
| missense   | NPFFR2     |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| NA         | NA         |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| upstream   | ENAM       |
| upstream   | AIM2       |
| missense   | CADM3      |
| intron     | CADM3      |
| missense   | ACKR1      |
| downstream | CADM3      |
| downstream | CADM3      |
| intergenic | CADM3-AS1  |
| intergenic | MPTX1      |
| intergenic | MPTX1      |
| intergenic | MPTX1      |
| intergenic | MPTX1      |
| intergenic | MPTX1      |
| upstream   | MPTX1      |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intergenic | AC112518.3 |
| intron     | RUFY3      |
| intron     | RUFY3      |
| intron     | RUFY3      |
| intron     | GRSF1      |

|               |               |
|---------------|---------------|
| intron        | GRSF1         |
| upstream      | RNU6-459P     |
| downstream    | RNU6-891P     |
| intergenic    | AC112518.3    |
| intergenic    | AC112518.3    |
| intron        | MOB1B         |
| intron        | PPEF2         |
| intron        | PPEF2         |
| intron        | HLA-B         |
| intron        | HLA-B         |
| intron        | FTO           |
| intron        | FTO           |
| upstream      | GSDMA         |
| intron        | GSDMA         |
| intron        | GSDMA         |
| intron        | GSDMA         |
| intron        | GSDMA         |
| intron        | GSDMA         |
| intron        | PSMD3         |
| downstream    | RP11-387H17.6 |
| intron        | MED24         |
| intron        | MED24         |
| splice_region | THRA          |
| intron        | THRA          |
| NA            | NA            |
| NA            | NA            |
| intron        | ZFPM2         |
| intron        | ZFPM2         |
| intron        | ZFPM2         |
| intron        | ZFPM2         |
| intron        | ZFPM2         |
| intron        | ZFPM2         |
| intron        | ZFPM2         |
| NA            | NA            |
| NA            | NA            |
| NA            | NA            |
| NA            | NA            |
| NA            | NA            |
| upstream      | MICB          |
| intron        | MICB          |
| NA            | NA            |
| intron        | MICB          |
| NA            | NA            |
| intron        | MICB          |
| intron        | MICB          |
| intron        | MICB          |

|            |               |
|------------|---------------|
| downstream | PPIAP9        |
| intron     | NFKBIL1       |
| intron     | NFKBIL1       |
| intron     | NFKBIL1       |
| downstream | LTB           |
| NA         | NA            |
| intron     | CFH           |
| intron     | CFH           |
| intron     | RP4-608O15.3  |
| intron     | RP4-608O15.3  |
| intron     | RP4-608O15.3  |
| intron     | RP4-608O15.3  |
| intron     | RP4-608O15.3  |
| intron     | RP4-608O15.3  |
| intron     | RP4-608O15.3  |
| intron     | CFHR4         |
| intron     | CFHR4         |
| intron     | RP4-608O15.3  |
| upstream   | CFHR5         |
| upstream   | CFHR5         |
| intron     | F13B          |
| intron     | CFH           |
| intron     | CFH           |
| intron     | CFH           |
| intron     | CFH           |
| intergenic | KCNT2         |
| intron     | KCNT2         |
| intron     | CFH           |
| synonymous | CFH           |
| intron     | CFH           |
| intron     | CFH           |
| intron     | CFH           |
| intron     | CFH           |
| intron     | CTD-2240J17.1 |
| intron     | CTD-2034I4.1  |
| intron     | IL16          |
| intron     | IL16          |
| intron     | HLA-DRB9      |
| intergenic | TBC1D22B      |
| upstream   | HLA-DQA1      |
| -          | -             |
| upstream   | RNU6-803P     |
| upstream   | RNU6-980P     |
| intron     | NLRP12        |
| intron     | NLRP12        |

|             |             |
|-------------|-------------|
| upstream    | NLRP12      |
| upstream    | NLRP12      |
| upstream    | NLRP12      |
| intergenic  | NLRP12      |
| intergenic  | NLRP12      |
| intergenic  | NLRP12      |
| intergenic  | NLRP12      |
| intergenic  | AC008753.6  |
| 3_prime_UTR | MYADM       |
| intron      | PRKCG       |
| intron      | CACNG7      |
| intron      | KCNT2       |
| intron      | CFH         |
| intron      | CFH         |
| intron      | CFH         |
| upstream    | RNU6-751P   |
| upstream    | AC008753.3  |
| upstream    | SEPT7P8     |
| upstream    | SEPT7P8     |
| upstream    | SEPT7P8     |
| upstream    | SEPT7P8     |
| downstream  | AC008753.3  |
| downstream  | AC008753.3  |
| intron      | NLRP12      |
| -           | -           |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| missense    | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| upstream    | NLRP12      |
| upstream    | NLRP12      |
| intergenic  | NLRP12      |
| intron      | CACNG7      |
| intron      | CTC-340D7.1 |
| intron      | CTC-340D7.1 |



|             |             |
|-------------|-------------|
| intron      | CTC-340D7.1 |
| upstream    | MIR525      |
| upstream    | MIR518F     |
| upstream    | MIR520C     |
| upstream    | MIR521-2    |
| upstream    | CTC-339O9.1 |
| upstream    | CTC-339O9.1 |
| upstream    | SEPT7P8     |
| upstream    | SEPT7P8     |
| upstream    | SEPT7P8     |
| upstream    | SEPT7P8     |
| downstream  | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| intron      | NLRP12      |
| missense    | NLRP12      |
| 5_prime_UTR | NLRP12      |
| upstream    | NLRP12      |
| upstream    | NLRP12      |
| upstream    | NLRP12      |
| intergenic  | NLRP12      |
| upstream    | FAM109A     |
| intron      | RP3-473L9.4 |
| upstream    | SRD5A2      |
| intergenic  | SRD5A2      |
| intergenic  | SRD5A2      |
| intergenic  | SRD5A2      |
| intergenic  | SRD5A2      |
| intergenic  | AL133247.3  |
| NA          | NA          |
| upstream    | COLEC11     |
| intron      | MASP1       |
| intron      | MASP1       |
| intron      | TSSC1       |
| intron      | TRAPPC12    |
| intron      | TRAPPC12    |
| intron      | TRAPPC12    |
| intron      | TRAPPC12    |
| intergenic  | AC018685.2  |
| intergenic  | AC011995.1  |

|                            |               |
|----------------------------|---------------|
| intergenic                 | AC011995.1    |
| intergenic                 | AC011995.1    |
| intergenic                 | AC011995.1    |
| intron                     | LINC01250     |
| intron                     | LINC01250     |
| intron                     | POLR1B        |
| intron                     | AC012442.5    |
| intron                     | AC012442.5    |
| intron                     | AC012442.5    |
| downstream                 | PSMD3         |
| downstream                 | RP11-387H17.6 |
| intron                     | CUX2          |
| 3_prime_UTR                | FAM109A       |
| intron                     | FAM109A       |
| intron                     | RP3-473L9.4   |
| intron                     | RP3-473L9.4   |
| intron                     | RP3-473L9.4   |
| intron                     | SH2B3         |
| intron                     | SH2B3         |
| intron                     | SH2B3         |
| intron                     | SH2B3         |
| 3_prime_UTR                | ATXN2         |
| intron                     | ATXN2         |
| intron                     | ATXN2         |
| intergenic                 | SLC20A1       |
| intergenic                 | NT5DC4        |
| missense                   | CKAP2L        |
| intergenic                 | IL1A          |
| downstream                 | IL1B          |
| non_coding_transcript_exon | RP11-67L14.1  |
| intergenic                 | XIAPP3        |
| intergenic                 | TBC1D22B      |
| intergenic                 | TBC1D22B      |
| intergenic                 | TBC1D22B      |
| intergenic                 | TBC1D22B      |
| intergenic                 | TBC1D22B      |
| intergenic                 | TBC1D22B      |
| intron                     | HLA-DRB5      |
| -                          | -             |
| NA                         | NA            |
| NA                         | NA            |
| NA                         | NA            |
| NA                         | NA            |
| intron                     | CFH           |
| intron                     | CFH           |
| NA                         | NA            |

|               |                   |
|---------------|-------------------|
| NA            | NA                |
| intron        | ZFPM2             |
| intron        | ZFPM2             |
| intron        | ZFPM2             |
| intron        | ZFPM2             |
| intron        | ZFPM2             |
| intron        | ZFPM2             |
| intron        | ZFPM2             |
| intron        | CFH               |
| intron        | CFH               |
| intron        | ZFPM2             |
| intron        | ZFPM2             |
| downstream    | NEU1              |
| downstream    | NEU1              |
| intron        | EHMT2             |
| missense      | EHMT2             |
| intron        | C2                |
| intron        | C2                |
| intron        | C2                |
| intron        | C2                |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| 3_prime_UTR   | NELFE             |
| 3_prime_UTR   | NELFE             |
| missense      | SKIV2L            |
| intron        | SKIV2L            |
| splice_region | DXO               |
| intron        | STK19             |
| intron        | STK19             |
| intron        | STK19             |
| intron        | STK19             |
| intron        | AL645922.1        |
| upstream      | AL645922.1        |
| upstream      | C4B               |
| intron        | C4B               |
| intron        | KCNT2             |
| intron        | KCNT2             |
| intron        | CFH               |
| intron        | CFH               |

|                            |               |
|----------------------------|---------------|
| intron                     | CFH           |
| intron                     | CFH           |
| intergenic                 | CFHR3         |
| 3_prime_UTR                | HLA-DPB1      |
| downstream                 | HLA-DPB2      |
| intergenic                 | AC112518.3    |
| intergenic                 | AC112518.3    |
| downstream                 | SLC25A48      |
| intron                     | GS1-39E22.1   |
| upstream                   | HLA-DRB1      |
| intron                     | HLA-DQA1      |
| intron                     | HLA-DQA1      |
| downstream                 | HLA-DQA1      |
| 5_prime_UTR                | SLFN11        |
| intergenic                 | RP11-686D22.6 |
| non_coding_transcript_exon | RP11-686D22.5 |
| intron                     | SLFN12L       |
| intron                     | SLFN12L       |
| intron                     | LZTFL1        |
| intron                     | FYCO1         |
| intron                     | LINC01405     |
| intergenic                 | LINC01405     |
| intron                     | NPFFR2        |
| intron                     | NPFFR2        |
| intergenic                 | MGAT5         |
| intergenic                 | MGAT5         |
| intergenic                 | MGAT5         |
| intergenic                 | MGAT5         |
| intergenic                 | MGAT5         |
| upstream                   | MGAT5         |
| intron                     | MGAT5         |
| intron                     | MGAT5         |
| intron                     | MGAT5         |
| upstream                   | TMEM102       |
| intron                     | ZBTB4         |
| intron                     | ZBTB4         |
| intron                     | ZBTB4         |
| synonymous                 | POLR2A        |
| intron                     | CUX2          |
| intron                     | CUX2          |
| intron                     | ALOX12-AS1    |
| intron                     | ALOX12        |
| intergenic                 | CTD-2545G14.2 |
| upstream                   | CTD-2545G14.2 |
| downstream                 | CTD-2545G14.2 |
| downstream                 | CTD-2545G14.2 |

|             |              |
|-------------|--------------|
| intron      | CLEC10A      |
| upstream    | CLEC10A      |
| intergenic  | CLEC10A      |
| downstream  | ASGR2        |
| downstream  | ASGR2        |
| -           | ASGR2        |
| intron      | ASGR2        |
| intron      | ASGR2        |
| intergenic  | ASGR2        |
| upstream    | RPL7AP64     |
| intergenic  | RPL7AP64     |
| intron      | SNX17        |
| intron      | PPM1G        |
| intergenic  | PPM1G        |
| intergenic  | NRBP1        |
| intergenic  | NRBP1        |
| intron      | IFT172       |
| downstream  | OR5K1        |
| intron      | CLDND1       |
| intron      | CPOX         |
| intergenic  | RNU6-461P    |
| upstream    | AC021660.1   |
| intergenic  | RNU6-461P    |
| intron      | PDE6A        |
| intron      | PDE6A        |
| intron      | SLC26A2      |
| intron      | HMGXB3       |
| intron      | HMGXB3       |
| intron      | HMGXB3       |
| intron      | HMGXB3       |
| synonymous  | HMGXB3       |
| intron      | HMGXB3       |
| intron      | HMGXB3       |
| intron      | HMGXB3       |
| -           | -            |
| intergenic  | RNU6-461P    |
| intergenic  | RNU6-461P    |
| intergenic  | CTC-281F24.2 |
| upstream    | RPL23AP73    |
| upstream    | RPL23AP73    |
| missense    | FBXO39       |
| intron      | TEKT1        |
| intron      | TEKT1        |
| intron      | TEKT1        |
| 3_prime_UTR | TEKT1        |
| intron      | TEKT1        |

|                            |              |
|----------------------------|--------------|
| intron                     | TEKT1        |
| intron                     | TEKT1        |
| upstream                   | TEKT1        |
| upstream                   | TEKT1        |
| intergenic                 | TEKT1        |
| intergenic                 | ALOX12P2     |
| intergenic                 | ALOX12P2     |
| intergenic                 | ALOX12P2     |
| upstream                   | ALOX12P2     |
| intron                     | ALOX12P2     |
| intron                     | ALOX12P2     |
| intron                     | ALOX12P2     |
| intron                     | ALOX12P2     |
| intron                     | ALOX12P2     |
| intron                     | ALOX12P2     |
| intron                     | ALOX12P2     |
| non_coding_transcript_exon | ALOX12P2     |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| intron                     | ALOX12-AS1   |
| non_coding_transcript_exon | RP11-530N7.2 |
| intron                     | ALOX12-AS1   |
| intron                     | CCR9         |
| intron                     | LZTFL1       |
| upstream                   | CCR2         |
| intron                     | LINC02009    |
| 5_prime_UTR                | AIM2         |
| intron                     | AIM2         |
| intron                     | AIM2         |
| intron                     | AIM2         |
| intron                     | AIM2         |
| intergenic                 | AIM2         |
| intron                     | CADM3        |
| intron                     | CADM3        |
| intron                     | CADM3        |
| missense                   | CADM3        |
| intron                     | CADM3        |
| missense                   | ACKR1        |
| downstream                 | CADM3        |

|            |               |
|------------|---------------|
| downstream | CADM3         |
| downstream | CADM3         |
| intergenic | CADM3-AS1     |
| intergenic | CADM3-AS1     |
| intergenic | CADM3-AS1     |
| intergenic | MPTX1         |
| -          | -             |
| intergenic | MPTX1         |
| downstream | LINC01565     |
| intron     | RPN1          |
| upstream   | RPN1          |
| intergenic | MRPL3         |
| intron     | CCDC13        |
| intron     | CCDC13        |
| intergenic | RP11-550P17.5 |
| intron     | RP11-550P17.5 |
| intron     | RP11-550P17.5 |
| intron     | RP11-180D21.3 |
| intron     | LZTFL1        |
| intron     | LZTFL1        |
| intron     | LZTFL1        |
| intron     | CCR9          |
| intron     | CCR9          |
| intergenic | LINC01989     |
| intergenic | CCL8          |
| intergenic | AC005549.3    |
| intergenic | CCL13         |
| intron     | ASIC2         |
| intron     | ASIC2         |
| intron     | ASIC2         |
| intron     | ASIC2         |
| intron     | ASIC2         |
| intron     | ASIC2         |
| -          | ASIC2         |
| intron     | ASIC2         |
| upstream   | ASIC2         |
| intergenic | AC084809.3    |
| intergenic | TMEM98        |
| intron     | SPACA3        |
| intron     | SPACA3        |
| intron     | ASIC2         |
| intron     | ASIC2         |
| upstream   | AIM2          |
| intron     | CADM3         |
| intron     | ACKR1         |
| missense   | ACKR1         |

|               |                   |
|---------------|-------------------|
| intergenic    | CADM3-AS1         |
| intergenic    | CADM3-AS1         |
| upstream      | HSPA1A            |
| intron        | SLC44A4           |
| intron        | SLC44A4           |
| intron        | SLC44A4           |
| intron        | SLC44A4           |
| upstream      | SLC44A4           |
| intron        | C2                |
| intron        | XXbac-BPG116M5.17 |
| splice_region | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| 3_prime_UTR   | NELFE             |
| 5_prime_UTR   | DXO               |
| intron        | STK19             |
| intron        | NAPRT             |
| intron        | EEF1D             |
| intergenic    | TSTA3             |
| intron        | ZNF623            |
| intron        | CCR3              |
| intergenic    | TKT               |
| intron        | HLA-DRA           |
| -             | HLA-DRB5          |
| intron        | SDAD1             |
| intron        | ART3              |
| intron        | ART3              |
| downstream    | NUP54             |
| -             | -                 |
| upstream      | MYL2              |
| upstream      | MYL2              |
| upstream      | MYL2              |
| upstream      | RP1-46F2.3        |
| intergenic    | LINC01405         |
| 3_prime_UTR   | SLFN5             |
| downstream    | SLFN12            |
| downstream    | BCAM              |
| intron        | NECTIN2           |
| missense      | APOE              |
| intron        | APOC4-APOC2       |
| intron        | CLPTM1            |
| downstream    | HLA-DRB6          |
| downstream    | HLA-DRB6          |
| upstream      | RNU1-61P          |
| intron        | HLA-DRB1          |
| intron        | HLA-DRB1          |
| upstream      | HLA-DRB1          |



|             |             |
|-------------|-------------|
| upstream    | HLA-DRB1    |
| intergenic  | TBC1D22B    |
| intergenic  | TBC1D22B    |
| intergenic  | TBC1D22B    |
| intergenic  | TBC1D22B    |
| intergenic  | TBC1D22B    |
| upstream    | HLA-DQA1    |
| intron      | HLA-DQA1    |
| intron      | HLA-DQA1    |
| intron      | HLA-DQA1    |
| -           | -           |
| missense    | HLA-DQA1    |
| intron      | HLA-DQB1    |
| intergenic  | TBC1D22B    |
| intergenic  | TBC1D22B    |
| downstream  | MTCO3P1     |
| upstream    | CCL18       |
| downstream  | CCL3        |
| upstream    | RP1-74B13.2 |
| intron      | RP1-74B13.2 |
| intron      | CCDC63      |
| intron      | CCDC63      |
| intron      | CCDC63      |
| intron      | MYL2        |
| upstream    | MYL2        |
| upstream    | MYL2        |
| intron      | RP1-46F2.3  |
| intron      | RP1-46F2.3  |
| upstream    | RP1-46F2.3  |
| intergenic  | LINC01405   |
| intergenic  | LINC01405   |
| intergenic  | LINC01405   |
| intergenic  | LINC01405   |
| upstream    | RASL10B     |
| intron      | GAS2L2      |
| upstream    | GAS2L2      |
| intron      | MMP28       |
| intron      | MMP28       |
| intron      | MMP28       |
| intron      | TAF15       |
| intron      | TAF15       |
| intron      | HEATR9      |
| intron      | HEATR9      |
| intron      | CCL5        |
| intron      | CCL5        |
| 3_prime_UTR | SLFN13      |

|             |               |
|-------------|---------------|
| downstream  | SLFN12L       |
| intergenic  | TKT           |
| intron      | CDCP1         |
| intron      | CDCP1         |
| downstream  | RPS24P8       |
| intergenic  | TKT           |
| upstream    | TLR10         |
| upstream    | TLR10         |
| missense    | TLR1          |
| intron      | TLR1          |
| intron      | TLR1          |
| upstream    | RP11-509I21.2 |
| upstream    | SNORD77       |
| upstream    | SNORD77       |
| upstream    | ALS2CL        |
| intron      | SLC6A20       |
| intron      | LZTFL1        |
| intergenic  | TKT           |
| intergenic  | TKT           |
| intergenic  | TKT           |
| intergenic  | TKT           |
| downstream  | FLT1P1        |
| intergenic  | TKT           |
| 3_prime_UTR | CCR1          |
| intron      | CCR3          |
| intron      | CCR3          |
| intron      | CCR3          |
| intron      | CCR3          |
| intergenic  | TKT           |
| intergenic  | TKT           |
| intron      | C17orf50      |
| intron      | HEATR9        |
| intron      | AP2B1         |
| intron      | AP2B1         |
| intron      | UNC45B        |
| intron      | UNC45B        |
| intron      | UNC45B        |
| intergenic  | SLC35G3       |
| intron      | RP11-799D4.4  |
| intergenic  | CTB-186H2.2   |
| upstream    | CCL23         |
| intron      | NFE2          |
| intron      | NFE2          |
| intron      | COPZ1         |
| intron      | COPZ1         |
| intron      | COPZ1         |

|             |               |
|-------------|---------------|
| intron      | COPZ1         |
| intron      | SERPINE2      |
| intron      | SERPINE2      |
| intron      | GALNT14       |
| intron      | CAPN14        |
| upstream    | CAPN14        |
| upstream    | EHD3          |
| intron      | EHD3          |
| -           | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| intron      | EHD3          |
| NA          | NA            |
| intron      | EHD3          |
| 3_prime_UTR | EHD3          |
| upstream    | AL121657.4    |
| downstream  | AL121657.4    |
| downstream  | AL121657.4    |
| intergenic  | XDH           |
| intergenic  | XDH           |
| downstream  | XDH           |
| NA          | NA            |
| NA          | NA            |
| intergenic  | RNU6-543P     |
| intergenic  | RNU6-543P     |
| intergenic  | RNU6-543P     |
| NA          | NA            |
| intergenic  | RNU6-543P     |
| intergenic  | RNU6-543P     |
| NA          | NA            |
| intergenic  | RP11-144G16.1 |
| downstream  | RP11-144G16.1 |
| downstream  | RP11-144G16.1 |
| upstream    | NRBF2         |
| NA          | NA            |
| intron      | JMJD1C        |
| intron      | JMJD1C        |
| NA          | NA            |
| intron      | JMJD1C        |

|            |                |
|------------|----------------|
| missense   | GP6            |
| intron     | GP6            |
| intron     | GP6            |
| intron     | RDH13          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| -          | -              |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| -          | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | ZFPM2          |
| intron     | PTPRC          |
| intron     | PTPRC          |
| downstream | RP11-686D22.4  |
| downstream | RP11-1094M14.7 |
| intron     | SLFN12L        |
| intron     | SLFN12L        |
| upstream   | SLFN12L        |
| missense   | SLFN14         |
| intron     | SLFN14         |
| intergenic | CCL1           |
| intergenic | AC011193.1     |
| intergenic | AC011193.1     |
| upstream   | AC011193.1     |
| intron     | SLFN12L        |
| downstream | SLFN14         |
| intergenic | RNU6-543P      |
| downstream | RP11-144G16.1  |

|                            |            |
|----------------------------|------------|
| upstream                   | GP6        |
| intron                     | RDH13      |
| NA                         | NA         |
| NA                         | NA         |
| NA                         | NA         |
| NA                         | NA         |
| NA                         | NA         |
| intron                     | TMC4       |
| missense                   | TMC4       |
| intron                     | MBOAT7     |
| intron                     | RPS9       |
| intron                     | RPS9       |
| downstream                 | LILRA6     |
| NA                         | NA         |
| intron                     | LILRA6     |
| NA                         | NA         |
| downstream                 | LILRB5     |
| upstream                   | LILRB5     |
| upstream                   | LILRB5     |
| -                          | AC010492.5 |
| non_coding_transcript_exon | RNU6-1307P |
| non_coding_transcript_exon | AC010492.5 |
| intron                     | AC010492.5 |
| intron                     | LILRB2     |
| intron                     | LILRB2     |
| intron                     | LILRB2     |
| intron                     | LILRB2     |
| missense                   | LILRB2     |
| upstream                   | LILRB2     |
| intergenic                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| downstream                 | AC010518.3 |
| upstream                   | AC010518.3 |
| upstream                   | AC010518.3 |
| upstream                   | AC010518.3 |
| upstream                   | AC010518.3 |
| intron                     | AC008984.5 |
| intron                     | AC008984.7 |

|             |               |
|-------------|---------------|
| downstream  | LILRA5        |
| downstream  | LILRA5        |
| downstream  | LILRA5        |
| downstream  | LILRA5        |
| downstream  | LILRA5        |
| downstream  | LILRA5        |
| intron      | LILRA5        |
| intron      | LILRA5        |
| upstream    | LILRA5        |
| upstream    | LILRA5        |
| intron      | AC008984.2    |
| downstream  | LILRA4        |
| -           | -             |
| intron      | LILRA4        |
| synonymous  | LILRA4        |
| downstream  | LAIR1         |
| NA          | NA            |
| NA          | NA            |
| missense    | MATN4         |
| NA          | NA            |
| downstream  | RBPJL         |
| downstream  | RBPJL         |
| 3_prime_UTR | SDC4          |
| intron      | SDC4          |
| intron      | SDC4          |
| intron      | SDC4          |
| intergenic  | RP11-323C15.1 |
| NA          | NA            |
| NA          | NA            |
| NA          | NA            |
| NA          | NA            |
| NA          | NA            |
| missense    | PIGT          |
| intron      | NECTIN2       |
| NA          | NA            |
| intron      | NECTIN2       |
| intron      | NECTIN2       |
| intron      | TOMM40        |
| synonymous  | TOMM40        |
| intron      | TOMM40        |
| upstream    | APOC1         |
| intron      | APOC1         |
| intron      | APOC1         |
| downstream  | APOC1         |
| downstream  | APOC1         |
| downstream  | APOC1         |

|              |             |
|--------------|-------------|
| upstream     | APOC1P1     |
| downstream   | APOC1P1     |
| upstream     | APOC4-APOC2 |
| intron       | APOC4-APOC2 |
| intron       | CLPTM1      |
| intron       | RELB        |
| intron       | RELB        |
| intergenic   | RNU6-461P   |
| intergenic   | RNU6-461P   |
| upstream     | AC021660.1  |
| intergenic   | RNU6-461P   |
| -            | AC021660.1  |
| intergenic   | RNU6-461P   |
| intergenic   | RNU6-461P   |
| intergenic   | RNU6-461P   |
| intergenic   | RNU6-461P   |
| splice_donor | ST3GAL6-AS1 |
| -            | ST3GAL6     |
| 3_prime_UTR  | PIGT        |
| intron       | WFDC2       |
| downstream   | SPINT3      |
| intron       | WFDC8       |
| intron       | WFDC8       |
| intron       | WFDC8       |
| intron       | NAE1        |
| intron       | PDP2        |
| intron       | PDP2        |
| intron       | CDH16       |
| intron       | RRAD        |
| intron       | CES2        |
| intron       | CES4A       |
| missense     | CES4A       |
| downstream   | CES4A       |
| upstream     | RN7SL543P   |
| intron       | CBFB        |
| intron       | CBFB        |
| intron       | CBFB        |
| NA           | NA          |
| NA           | NA          |
| NA           | NA          |
| NA           | NA          |
| NA           | NA          |
| intergenic   | ABO         |
| NA           | NA          |
| NA           | NA          |
| downstream   | ABO         |

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|            |               |
|------------|---------------|
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| -          | -             |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| intron     | RAB11FIP3     |
| downstream | MED25         |
| intron     | PTOV1         |
| intron     | IL4I1         |
| intron     | IL4I1         |
| intron     | ATF5          |
| upstream   | SIGLEC11      |
| downstream | VRK3          |
| downstream | VRK3          |
| downstream | VRK3          |
| intron     | VRK3          |
| intron     | VRK3          |
| intron     | MICB          |
| upstream   | TNF           |
| intron     | SYT3          |
| intron     | C19orf81      |
| intron     | C19orf81      |
| intron     | C19orf81      |
| intron     | SHANK1        |
| intron     | SHANK1        |
| upstream   | SHANK1        |
| downstream | CLEC11A       |
| intergenic | CLEC11A       |
| intergenic | CLEC11A       |
| intron     | CTD-2568A17.1 |
| intron     | ACPT          |
| intron     | CHST11        |
| intron     | CHST11        |
| intron     | CHST11        |

|            |               |
|------------|---------------|
| intron     | CHST11        |
| intron     | CHST11        |
| intron     | CHST11        |
| intron     | CHST11        |
| intergenic | TKT           |
| intron     | TMIE          |
| intron     | DUS2          |
| intron     | DUS2          |
| intron     | MSI2          |
| intron     | MSI2          |
| intron     | MSI2          |
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| intron     | MSI2          |
| intron     | MSI2          |
| intron     | MYH14         |
| intron     | MYH14         |
| intron     | MYH14         |
| intron     | MYH14         |
| intron     | RBBP5         |
| intron     | DSTYK         |
| intergenic | RP11-626I20.2 |
| intergenic | RP11-626I20.3 |
| upstream   | RP11-626I20.3 |
| upstream   | RP11-626I20.3 |
| intergenic | RP11-626I20.3 |
| intergenic | STAB2         |
| intergenic | STAB2         |

|               |              |
|---------------|--------------|
| intron        | STAB2        |
| intron        | STAB2        |
| intron        | STAB2        |
| downstream    | NT5DC3       |
| intron        | MYH14        |
| intron        | KCNC3        |
| intron        | KCNC3        |
| intron        | NR1H2        |
| intron        | NR1H2        |
| intron        | NR1H2        |
| intron        | NR1H2        |
| intron        | CTD-2545M3.6 |
| intron        | CTD-2545M3.6 |
| intron        | CTD-2545M3.6 |
| missense      | MYBPC2       |
| intron        | MYBPC2       |
| intron        | MYBPC2       |
| intron        | MYBPC2       |
| intron        | FAM71E1      |
| intron        | FAM71E1      |
| intron        | FAM71E1      |
| intron        | FAM71E1      |
| synonymous    | EMC10        |
| -             | -            |
| 3_prime_UTR   | EMC10        |
| 3_prime_UTR   | EMC10        |
| downstream    | EMC10        |
| downstream    | EMC10        |
| downstream    | EMC10        |
| intron        | ABCA6        |
| intron        | ABCA10       |
| intron        | ABCA10       |
| splice_region | ABCA5        |
| intron        | ABCA5        |
| NA            | NA           |
| NA            | NA           |
| intron        | CUX2         |
| NA            | NA           |
| intron        | CUX2         |
| intron        | CUX2         |
| NA            | NA           |
| intron        | RP3-473L9.4  |
| intron        | SH2B3        |
| intron        | ATXN2        |
| intron        | ATXN2        |
| intron        | ATXN2        |

|             |              |
|-------------|--------------|
| intron      | ATXN2        |
| intron      | ATXN2        |
| NA          | NA           |
| NA          | NA           |
| NA          | NA           |
| NA          | NA           |
| NA          | NA           |
| NA          | NA           |
| NA          | NA           |
| intron      | NRBF2        |
| NA          | NA           |
| intron      | JMJD1C       |
| intron      | JMJD1C       |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| intron      | CFH          |
| 5_prime_UTR | CFHR3        |
| 3_prime_UTR | RP13-512J5.1 |
| NA          | NA           |
| intron      | RP13-512J5.1 |
| NA          | NA           |
| NA          | NA           |
| upstream    | RNASEH1      |
| intergenic  | RNASEH1-AS1  |
| NA          | NA           |
| intergenic  | RPS7         |
| intergenic  | COLEC11      |
| upstream    | COLEC11      |
| NA          | NA           |
| NA          | NA           |
| intron      | COLEC11      |
| intron      | COLEC11      |
| intron      | COLEC11      |
| intron      | COLEC11      |
| NA          | NA           |
| downstream  | COLEC11      |
| intron      | ALLC         |
| intron      | ALLC         |

|             |               |
|-------------|---------------|
| NA          | NA            |
| intron      | ALLC          |
| NA          | NA            |
| NA          | NA            |
| NA          | NA            |
| NA          | NA            |
| intergenic  | DSG1          |
| intergenic  | DSG1          |
| NA          | NA            |
| NA          | NA            |
| intergenic  | DSG1          |
| NA          | NA            |
| intergenic  | DSG1          |
| intron      | KCNT2         |
| intron      | CFH           |
| intron      | CFH           |
| intron      | CFH           |
| intron      | CFH           |
| intron      | CFH           |
| intron      | CFH           |
| intron      | DSG4          |
| 3_prime_UTR | DSG3          |
| intron      | PHLDB3        |
| intron      | PHLDB3        |
| intron      | XRCC1         |
| missense    | PINLYP        |
| downstream  | PLAUR         |
| missense    | PLAUR         |
| intron      | PLAUR         |
| NA          | NA            |
| NA          | NA            |
| intergenic  | RP11-675P14.1 |
| intergenic  | RP11-675P14.1 |
| intergenic  | RP11-675P14.1 |
| NA          | NA            |
| NA          | NA            |
| intergenic  | RP11-675P14.1 |
| intron      | RP13-512J5.1  |
| intron      | RP13-512J5.1  |
| intron      | RP13-512J5.1  |
| intron      | RP13-512J5.1  |
| intron      | RP13-512J5.1  |
| intron      | RP13-512J5.1  |
| intron      | RNASEH1       |
| synonymous  | RNASEH1       |
| intron      | RNASEH1       |

[illegible]

|               |                   |
|---------------|-------------------|
| NA            | NA                |
| intron        | ALLC              |
| intron        | ALLC              |
| NA            | NA                |
| intron        | ALLC              |
| NA            | NA                |
| 5_prime_UTR   | C6orf48           |
| intron        | SLC44A4           |
| intron        | C2                |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | XXbac-BPG116M5.17 |
| intron        | NELFE             |
| 5_prime_UTR   | NELFE             |
| splice_region | STK19             |
| intron        | VLDLR-AS1         |
| upstream      | VLDLR             |
| intergenic    | VLDLR             |
| intergenic    | VLDLR             |
| -             | -                 |
| intron        | PUM3              |
| downstream    | RP1-261G23.5      |
| intron        | RP1-261G23.5      |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
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| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intron        | RP5-1120P11.1     |
| intron        | RP5-1120P11.1     |
| intron        | RP5-1120P11.1     |
| upstream      | CRIP3             |
| 3_prime_UTR   | ZNF318            |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |
| intergenic    | ACTG1P9           |

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|            |               |
|------------|---------------|
| intergenic | ACTG1P9       |
| upstream   | C6orf223      |
| upstream   | C6orf223      |
| downstream | C6orf223      |
| intron     | RP5-1120P11.1 |
| intron     | RP5-1120P11.1 |
| intron     | RP5-1120P11.1 |
| intron     | RP5-1120P11.1 |
| intron     | RP5-1120P11.1 |
| intron     | RP11-344J7.3  |
| intron     | RP11-344J7.3  |
| intergenic | ACTG1P9       |
| intergenic | ACTG1P9       |
| intergenic | ACTG1P9       |
| intergenic | ACTG1P9       |
| intergenic | ACTG1P9       |
| intergenic | EGR2          |
| intergenic | RNU6-543P     |
| intergenic | RNU6-543P     |
| intergenic | RNU6-543P     |
| intergenic | RNU6-543P     |
| -          | RNU6-543P     |
| downstream | RNU6-543P     |
| intergenic | RNU6-543P     |
| intergenic | RP11-144G16.1 |
| downstream | RP11-144G16.1 |
| downstream | RP11-144G16.1 |
| upstream   | RP11-144G16.1 |
| upstream   | NRBF2         |
| upstream   | NRBF2         |
| intron     | NRBF2         |
| intron     | NRBF2         |
| downstream | NRBF2         |

## **Cytokine**

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