Stroke SNP Documentation using LaTeX

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> setwd("/home/pouria3/workspace/GenStat/trunk/Hyp")
> source("Snp2.R")
Found a statistically insignificant OR :
       PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus ORValue     CI FreqControl FreqPop ModelType Pvalue
22 19752551 rs3900940      Caucasian          Vienna      CT    1.26 1-1.59       0.406      NA  additive   0.05
Found a statistically insignificant OR :
       PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus ORValue       CI FreqControl FreqPop ModelType Pvalue
23 19752551 rs3900940      Caucasian          Vienna      CC    1.19 0.8-1.75       0.092      NA  additive   0.25
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in Genotype sec;quering SNPInfo for frequencies ...
      PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus ORValue        CI FreqControl FreqPop ModelType Pvalue
1 19403135 rs1062708          Asian        Japanese      TT    1.00      <NA>        21.7      NA recessive     NA
2 19403135 rs1062708          Asian        Japanese      CC    0.82 0.68-0.99        29.6      NA recessive 0.0374
3 19403135 rs1062708          Asian        Japanese      TC    1.00      <NA>        48.7      NA recessive     NA
[1] "Frequencies are given, I am not going to hapmap!"
[1] "Frequencies are given, I am not going to hapmap!"
[1] "Frequencies are given, I am not going to hapmap!"
Data Gathering Completed, Solving the equations ...
solving non linear equations ...
 Freqs Are:  [1] 48.7 21.7 29.6
  Successful convergence.
1.843298e-05 1.844085e-05 1.512154e-05
      snpID Genotype freq   OR       NewOR BroadEthnicity NarrowEthnicity   name
1 rs1062708     <NA>  0.0 0.00 0.000000000          Asian        Japanese RUVBL2
2 rs1062708       TC 48.7 1.00 0.010542391          Asian        Japanese RUVBL2
3 rs1062708       TT 21.7 1.00 0.010546894          Asian        Japanese RUVBL2
4 rs1062708       CC 29.6 0.82 0.008648451          Asian        Japanese RUVBL2
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in Allele sec;quering SNPInfo for Major Minor Allele ...
       PMID      dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Freq
4 19369658 rs10794579      Caucasian White_Framingham Heart Study_Rotterdam Study       T    1.24 1.13-1.35          NA    0
5 19369658 rs10794579      Caucasian White_Framingham Heart Study_Rotterdam Study       C    1.00      <NA>          NA    0
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      TT    Het     CC
1 0.1849 0.4902 0.3249
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.3249 0.4902 0.1849
  Successful convergence.
0.001434654 0.001778309 0.002204374
       snpID Genotype   freq     OR     NewOR BroadEthnicity                                NarrowEthnicity     name
1 rs10794579     <NA> 0.0000 0.0000 0.0000000      Caucasian White_Framingham Heart Study_Rotterdam Study C10orf88
2 rs10794579       CC 0.3249 1.0000 0.8216869      Caucasian White_Framingham Heart Study_Rotterdam Study C10orf88
3 rs10794579       TC 0.4902 1.2400 1.0188633      Caucasian White_Framingham Heart Study_Rotterdam Study C10orf88
4 rs10794579       TT 0.1849 1.5376 1.2635116      Caucasian White_Framingham Heart Study_Rotterdam Study C10orf88
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in Allele sec;quering SNPInfo for Major Minor Allele ...
       PMID      dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Freq
6 19369658 rs10837576      Caucasian White_Framingham Heart Study_Rotterdam Study       G    1.00      <NA>          NA    0
7 19369658 rs10837576      Caucasian White_Framingham Heart Study_Rotterdam Study       A    0.78 0.70-0.85          NA    0
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      GG    Het     AA
1 0.5041 0.4118 0.0841
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.5041 0.4118 0.0841
  Successful convergence.
0.001991132 0.001553653 0.00121235
       snpID Genotype   freq     OR     NewOR BroadEthnicity                                NarrowEthnicity           name
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1 rs10837576     <NA> 0.0000 0.0000 0.0000000      Caucasian White_Framingham Heart Study_Rotterdam Study LRRC4C_RPL9P23
2 rs10837576       GG 0.5041 1.0000 1.1410412      Caucasian White_Framingham Heart Study_Rotterdam Study LRRC4C_RPL9P23
3 rs10837576       GA 0.4118 0.7800 0.8899488      Caucasian White_Framingham Heart Study_Rotterdam Study LRRC4C_RPL9P23
4 rs10837576       AA 0.0841 0.6084 0.6942094      Caucasian White_Framingham Heart Study_Rotterdam Study LRRC4C_RPL9P23
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in Allele sec;quering SNPInfo for Major Minor Allele ...
       PMID      dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Freq
8 19369658 rs11615969      Caucasian White_Framingham Heart Study_Rotterdam Study       T    1.00      <NA>          NA    0
9 19369658 rs11615969      Caucasian White_Framingham Heart Study_Rotterdam Study       C    1.53 1.27-1.84          NA    0
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      TT    Het     CC
1 0.8281 0.1638 0.0081
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.8281 0.1638 0.0081
  Successful convergence.
0.001590499 0.002431315 0.003715138
       snpID Genotype   freq     OR     NewOR BroadEthnicity                                NarrowEthnicity            name
1 rs11615969     <NA> 0.0000 0.0000 0.0000000      Caucasian White_Framingham Heart Study_Rotterdam Study BBS10_LOC641695
2 rs11615969       TT 0.8281 1.0000 0.9110882      Caucasian White_Framingham Heart Study_Rotterdam Study BBS10_LOC641695
3 rs11615969       TC 0.1638 1.5300 1.3939080      Caucasian White_Framingham Heart Study_Rotterdam Study BBS10_LOC641695
4 rs11615969       CC 0.0081 2.3409 2.1326867      Caucasian White_Framingham Heart Study_Rotterdam Study BBS10_LOC641695
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID      dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Fre
10 19369658 rs12786704      Caucasian White_Framingham Heart Study_Rotterdam Study       G    1.36 1.19-1.55          NA    (
11 19369658 rs12786704      Caucasian White_Framingham Heart Study_Rotterdam Study       A    1.00      <NA>          NA    (
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      GG    Het     AA
1 0.0256 0.2688 0.7056
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.7056 0.2688 0.0256
  Successful convergence.
0.001560862 0.00212162 0.002883
       snpID Genotype   freq     OR     NewOR BroadEthnicity                                NarrowEthnicity name
1 rs12786704     <NA> 0.0000 0.0000 0.0000000      Caucasian White_Framingham Heart Study_Rotterdam Study  NTM
2 rs12786704       AA 0.7056 1.0000 0.8940847      Caucasian White_Framingham Heart Study_Rotterdam Study  NTM
3 rs12786704       AG 0.2688 1.3600 1.2159778      Caucasian White_Framingham Heart Study_Rotterdam Study  NTM
4 rs12786704       GG 0.0256 1.8496 1.6536139      Caucasian White_Framingham Heart Study_Rotterdam Study  NTM
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in Genotype sec;quering SNPInfo for frequencies ...
       PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus ORValue        CI FreqControl FreqPop ModelType Pvalue
12 19403135 rs1671021          Asian        Japanese      CT    1.00      <NA>        24.9      NA  dominant     NA
13 19403135 rs1671021          Asian        Japanese      CC    0.74 0.62-0.87         2.2      NA  dominant  0.004
14 19403135 rs1671021          Asian        Japanese      TT    1.00      <NA>        72.9      NA  dominant     NA
[1] "Frequencies are given, I am not going to hapmap!"
[1] "Frequencies are given, I am not going to hapmap!"
[1] "Frequencies are given, I am not going to hapmap!"
Data Gathering Completed, Solving the equations ...
solving non linear equations ...
 Freqs Are:  [1] 72.9 24.9  2.2
  Successful convergence.
1.755488e-05 1.755475e-05 1.299052e-05
      snpID Genotype freq   OR       NewOR BroadEthnicity NarrowEthnicity  name
1 rs1671021     <NA>  0.0 0.00 0.000000000          Asian        Japanese LLGL2
2 rs1671021       TT 72.9 1.00 0.010040170          Asian        Japanese LLGL2
3 rs1671021       CT 24.9 1.00 0.010040098          Asian        Japanese LLGL2
4 rs1671021       CC  2.2 0.74 0.007429639          Asian        Japanese LLGL2
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus ORValue   CI FreqControl FreqPop ModelType Pvalue
15 21511255 rs1671021      Caucasian      Portuguese       C 1.00001 <NA>          NA      NA  Additive  0.965
16 21511255 rs1671021      Caucasian      Portuguese       T 1.00000 <NA>          NA      NA  Additive     NA
Calculate Genotype Frequencies based on Hardy-Weinberg principle
No Freq Found... Going to HapMap
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.4159292 0.4247788 0.1592920
  Successful convergence.
0.001745407 0.001745519 0.001745528
      snpID Genotype      freq      OR     NewOR BroadEthnicity NarrowEthnicity  name
1 rs1671021     <NA> 0.0000000 0.00000 0.0000000      Caucasian      Portuguese LLGL2
2 rs1671021       TT 0.4159292 1.00000 0.9999795      Caucasian      Portuguese LLGL2
3 rs1671021       TC 0.4247788 1.00001 1.0000438      Caucasian      Portuguese LLGL2
4 rs1671021       CC 0.1592920 1.00002 1.0000486      Caucasian      Portuguese LLGL2
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID      dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Fre
17 19369658 rs17429019      Caucasian White_Framingham Heart Study_Rotterdam Study       A    1.00      <NA>          NA    0
18 19369658 rs17429019      Caucasian White_Framingham Heart Study_Rotterdam Study       G    1.49 1.26-1.76          NA    0
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      AA    Het     GG
1 0.8836 0.1128 0.0036
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.8836 0.1128 0.0036
  Successful convergence.
0.001647399 0.002452685 0.003650139
       snpID Genotype   freq     OR    NewOR BroadEthnicity                                NarrowEthnicity       name
1 rs17429019     <NA> 0.0000 0.0000 0.000000      Caucasian White_Framingham Heart Study_Rotterdam Study TPRG1_TP63
2 rs17429019       AA 0.8836 1.0000 0.943736      Caucasian White_Framingham Heart Study_Rotterdam Study TPRG1_TP63
3 rs17429019       AG 0.1128 1.4900 1.406190      Caucasian White_Framingham Heart Study_Rotterdam Study TPRG1_TP63
4 rs17429019       GG 0.0036 2.2201 2.095237      Caucasian White_Framingham Heart Study_Rotterdam Study TPRG1_TP63
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID     dbSNP BroadEthnicity                               NarrowEthnicity AG_Plus ORValue        CI FreqControl Freq
19 19369658 rs2318308      Caucasian White_Framingham Heart Study_Rotterdam Study       G    1.00      <NA>          NA    0
20 19369658 rs2318308      Caucasian White_Framingham Heart Study_Rotterdam Study       A    0.75 0.66-0.85          NA    0
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      GG    Het     AA
1 0.5329 0.3942 0.0729
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.5329 0.3942 0.0729
  Successful convergence.
0.002006844 0.00150589 0.0011298
      snpID Genotype   freq     OR     NewOR BroadEthnicity                                NarrowEthnicity          name
1 rs2318308     <NA> 0.0000 0.0000 0.0000000      Caucasian White_Framingham Heart Study_Rotterdam Study FTHL13_YBX1P1
2 rs2318308       GG 0.5329 1.0000 1.1500633      Caucasian White_Framingham Heart Study_Rotterdam Study FTHL13_YBX1P1
3 rs2318308       GA 0.3942 0.7500 0.8625483      Caucasian White_Framingham Heart Study_Rotterdam Study FTHL13_YBX1P1
4 rs2318308       AA 0.0729 0.5625 0.6468867      Caucasian White_Framingham Heart Study_Rotterdam Study FTHL13_YBX1P1
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in Genotype sec;quering SNPInfo for frequencies ...
       PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus  ORValue       CI FreqControl FreqPop ModelType Pvalue
21 19752551 rs3900940      Caucasian          Vienna      TT 1.000000     <NA>       0.503      NA  additive     NA
22 19752551 rs3900940      Caucasian          Vienna      CT 1.000001   1-1.59       0.406      NA  additive   0.05
23 19752551 rs3900940      Caucasian          Vienna      CC 1.000001 0.8-1.75       0.092      NA  additive   0.25
[1] "Frequencies are given, I am not going to hapmap!"
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[1] "Frequencies are given, I am not going to hapmap!"
[1] "Frequencies are given, I am not going to hapmap!"
Data Gathering Completed, Solving the equations ...
solving non linear equations ...
 Freqs Are:  [1] 0.092 0.406 0.503
  Successful convergence.
0.001743675 0.001743759 0.001743619
      snpID Genotype  freq       OR     NewOR BroadEthnicity NarrowEthnicity  name
1 rs3900940     <NA> 0.000 0.000000 0.0000000      Caucasian          Vienna MYH15
2 rs3900940       CC 0.092 1.000001 0.9989853      Caucasian          Vienna MYH15
3 rs3900940       CT 0.406 1.000001 0.9990338      Caucasian          Vienna MYH15
4 rs3900940       TT 0.503 1.000000 0.9989532      Caucasian          Vienna MYH15
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID     dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Freq
24 19369658 rs4867131      Caucasian White_Framingham Heart Study_Rotterdam Study       A    1.49 1.27-1.75          NA    0
25 19369658 rs4867131      Caucasian White_Framingham Heart Study_Rotterdam Study       C    1.00      <NA>          NA    0
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      AA    Het     CC
1 0.0121 0.1958 0.7921
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.7921 0.1958 0.0121
  Successful convergence.
0.001571832 0.00234025 0.003482978
      snpID Genotype   freq     OR    NewOR BroadEthnicity                                NarrowEthnicity            name
1 rs4867131     <NA> 0.0000 0.0000 0.000000      Caucasian White_Framingham Heart Study_Rotterdam Study LOC340113_RPS8P8
2 rs4867131       CC 0.7921 1.0000 0.900378      Caucasian White_Framingham Heart Study_Rotterdam Study LOC340113_RPS8P8
3 rs4867131       CA 0.1958 1.4900 1.341576      Caucasian White_Framingham Heart Study_Rotterdam Study LOC340113_RPS8P8
4 rs4867131       AA 0.0121 2.2201 1.998949      Caucasian White_Framingham Heart Study_Rotterdam Study LOC340113_RPS8P8
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in Genotype sec;quering SNPInfo for frequencies ...
       PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus ORValue        CI FreqControl FreqPop ModelType Pvalue
26 19403135 rs6007897          Asian        Japanese      TT    1.00      <NA>        97.3      NA  dominant     NA
27 19403135 rs6007897          Asian        Japanese      CC    1.85 1.29-2.61         0.0      NA  dominant  6e-04
28 19403135 rs6007897          Asian        Japanese      CT    1.85 1.29-2.61         2.7      NA  dominant  6e-04
[1] "Frequencies are given, I am not going to hapmap!"
[1] "Frequencies are given, I am not going to hapmap!"
[1] "Frequencies are given, I am not going to hapmap!"
Data Gathering Completed, Solving the equations ...
solving non linear equations ...
 Freqs Are:  [1] 97.3  0.0  2.7
  Successful convergence.
1.70635e-05 3.154798e-05 3.154227e-05
      snpID Genotype freq   OR       NewOR BroadEthnicity NarrowEthnicity   name
1 rs6007897     <NA>  0.0 0.00 0.000000000          Asian        Japanese CELSR1
2 rs6007897       TT 97.3 1.00 0.009759129          Asian        Japanese CELSR1
3 rs6007897       CC  0.0 1.85 0.018043502          Asian        Japanese CELSR1
4 rs6007897       CT  2.7 1.85 0.018040240          Asian        Japanese CELSR1
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID     dbSNP BroadEthnicity NarrowEthnicity AG_Plus ORValue        CI FreqControl FreqPop ModelType Pvalue
29 21511255 rs6007897      Caucasian      Portuguese       C    1.43 1.13-1.81          NA      NA  Additive  0.003
30 21511255 rs6007897      Caucasian      Portuguese       T    1.00      <NA>          NA      NA  Additive     NA
Calculate Genotype Frequencies based on Hardy-Weinberg principle
No Freq Found... Going to HapMap
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.66666667 0.31531532 0.01801802
  Successful convergence.
0.001512433 0.002161409 0.003087972
      snpID Genotype       freq     OR     NewOR BroadEthnicity NarrowEthnicity   name
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1 rs6007897     <NA> 0.00000000 0.0000 0.0000000      Caucasian      Portuguese CELSR1
2 rs6007897       TT 0.66666667 1.0000 0.8663014      Caucasian      Portuguese CELSR1
3 rs6007897       TC 0.31531532 1.4300 1.2388318      Caucasian      Portuguese CELSR1
4 rs6007897       CC 0.01801802 2.0449 1.7715446      Caucasian      Portuguese CELSR1
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID     dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Freq
31 19369658 rs6820391      Caucasian White_Framingham Heart Study_Rotterdam Study       C    1.00      <NA>          NA    0
32 19369658 rs6820391      Caucasian White_Framingham Heart Study_Rotterdam Study       A    1.24 1.13-1.36          NA    0
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      CC    Het     AA
1 0.5184 0.4032 0.0784
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.5184 0.4032 0.0784
  Successful convergence.
0.001532941 0.00190014 0.002355106
      snpID Genotype   freq     OR     NewOR BroadEthnicity                                NarrowEthnicity name
1 rs6820391     <NA> 0.0000 0.0000 0.0000000      Caucasian White_Framingham Heart Study_Rotterdam Study LNX1
2 rs6820391       CC 0.5184 1.0000 0.8780664      Caucasian White_Framingham Heart Study_Rotterdam Study LNX1
3 rs6820391       CA 0.4032 1.2400 1.0887978      Caucasian White_Framingham Heart Study_Rotterdam Study LNX1
4 rs6820391       AA 0.0784 1.5376 1.3501132      Caucasian White_Framingham Heart Study_Rotterdam Study LNX1
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in Allele sec;quering SNPInfo for Major Minor Allele ...
        PMID    dbSNP BroadEthnicity                                NarrowEthnicity AG_Plus ORValue        CI FreqControl Freql
33 19369658 rs713536      Caucasian White_Framingham Heart Study_Rotterdam Study       T    1.26 1.14-1.38          NA    0.4
34 19369658 rs713536      Caucasian White_Framingham Heart Study_Rotterdam Study       C    1.00      <NA>          NA    0.5
Calculate Genotype Frequencies based on Hardy-Weinberg principle
Using FreqPop ...
Freqencies are :
      TT    Het     CC
1 0.2209 0.4982 0.2809
[1] "Data Gathering Completed, Solving the equations ..."
solving non linear equations ...
 Freqs Are:  [1] 0.2809 0.4982 0.2209
  Successful convergence.
0.00138647 0.001746381 0.002199435
     snpID Genotype   freq     OR     NewOR BroadEthnicity                                NarrowEthnicity         name
1 rs713536     <NA> 0.0000 0.0000 0.0000000      Caucasian White_Framingham Heart Study_Rotterdam Study C8orf79_DLC1
2 rs713536       CC 0.2809 1.0000 0.7940517      Caucasian White_Framingham Heart Study_Rotterdam Study C8orf79_DLC1
3 rs713536       CT 0.4982 1.2600 1.0005384      Caucasian White_Framingham Heart Study_Rotterdam Study C8orf79_DLC1
4 rs713536       TT 0.2209 1.5876 1.2606747      Caucasian White_Framingham Heart Study_Rotterdam Study C8orf79_DLC1
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Number of Snps processed is :  13
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