Genome.wide	chromosome.5.inversion	Chromosome.8.small.inversion	Chromosome.8.inversion..large.inversion.including.small.inversion.	Chromosome.8.inversion..large.inversion.without.small.inversion.	Chromosome.inversion.14
L1 nonsynonymous SNV	0	41	802	761	0
unknown	0	0	0	0	0
nonframeshift insertion	0	15	202	187	0
stop gain	0	9	98	89	0
frameshift deletion	0	11	211	200	0

Genome. wide	chromosome.5. inversion	Chromosome.8.small.inversion	Chromosome.8.inversion..large.inversion.including.small.inversion.	Chromosome.8.inversion..large.inversion. without.small.inversion.	Chromosome.inversion.14
nonframe shift deletion	0	10	196	186	0
frameshift insertion	0	12	195	183	0
stop loss	0	4	35	31	0
intronic	0	37	625	588	0
UTR3	0	19	318	299	0
UTR5	0	20	261	241	0
splicing	0	9	146	137	0

S1

Genome.wide	chromosome.5.inversion	Chromosome.8.small.inversion	Chromosome.8.inversion..large.inversion.including.small.inversion.	Chromosome.8.inversion..large.inversion.without.small.inversion.	Chromosome.inversion.14
nonsynonymous SNV	0	0	0	0	0
unknown	0	0	0	0	0
nonframeshift insertion	0	0	0	0	0
stop gain	0	0	0	0	0
frameshift deletion	0	0	0	0	0
nonframeshift deletion	0	0	0	0	0

Genome. wide	chromosome.5. inversion	Chromosome.8.small.inversion	Chromosome.8.inversion..large.inversion.including.small.inversion.	Chromosome.8.inversion..large.inversion.without.small.inversion.	Chromosome.inversion.14
frameshift insertion	0	0	0	0	0
stop loss	0	0	0	0	0
intronic	0	0	0	0	0
UTR3	0	0	0	0	0
UTR5	0	0	0	0	0
splicing	0	0	0	0	0