

	symbol	Chr.name	start.pos	symb.LOC	entrez
1	0	5	45	168	0
2	0	5	44	171	0
3	0	6	44	167	0
4	0	4	44	165	0

Table 1: All columns except symb.Loc show the number of missing or NA values. symb.Loc shows the number of genes symbols that still begin with LOC.