24,820 sequences containing at least 40bp non-Alu sequences

8,738 sequences unambiguously mapped to \textit{in silico} bisulfite converted reference genome

1,762 sequences (327 clusters) with a putative insertion site at least 10 bp from the most adjacent Alu

316 clusters identified as putative novel Alu integrations (not present in either dbSNP or dbRIP)