GENOME WIDE ANALYSIS OF THE HUMAN ALU YB-LINEAGE

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The Alu Yb-lineage is a "young" largely human-specific group of SINE (short interspersed element) subfamilies that have integrated throughout the human genome. In this study, we have computationally analyzed the draft sequence of the human genome for Alu Yb-lineage subfamily members present on autosomal chromosomes. A total of 1733 Yb Alu subfamily members have integrated into human autosomes. The estimated average age of Yb-lineage subfamilies, Yb7, Yb8 and Yb9, was estimated as 4.81, 2.39 and 2.32 million years, respectively. In order to determine the contribution of the Alu Yb lineage to human genomic diversity, 1202 polymerase chain reaction (PCR)-based assays were performed using oligonucleotide primers that amplified the genomic regions containing individual Yblineage subfamily members. Approximately 20% of the *Alu* elements were polymorphic for insertion presence/absence in the human genome. Fewer than 0.5% of the Yb Alus were also present at orthologous positions in non-human primate genomes. Sequence analysis of these loci showed that each of the orthologous loci from non-human primate genomes contained Y, Sg and Sx Alu family members. Thus, these Yb Alu elements are thought to have integrated into the human genome guite recently and replaced the older Alu elements via one of three mechanisms: gene conversion, Alumediated deletion or parallel, independent insertion. The high copy number, level of insertion polymorphism, and estimated average age suggests that many members of the Alu Yb elements will be useful in a wide range of genetic analyses.