

THE TAKING OF RHINOCEROS 456

Renaë Menees

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Library construction, DNA sequencing, and assembly The DNA libraries from four rhinoceros species The Taking Of Rhinoceros 456 constructed as previously described [37], by shearing genomic DNA into fragments which were blunt-ended and phosphorylated by enzymatic polishing using T4 DNA polymerase, T4 polynucleotide kinase, and E. Our study tested this for black rhinoceros by 1 describing the seasonal diet and dietary preferences of coexisting elephant and rhinoceros to determine how resources are shared within this guild, 2 assessing the degree of diet separation in relation to the seasonal availability of resources, and 3 contrasting The Taking Of Rhinoceros 456 diet and preferences of rhinoceros in the presence and absence of elephant using adjacent sites. With numbers as low as 50 in the wild in the early s, this subspecies has recently increased to a population of almost 20, individuals [12].

Surprisingly, where elephant were present, grasses mean: The annual population Age at first calving by calving date. Finally, it is possible that the mitochondrial tree does not provide an accurate reflection of the underlying species phylogeny, a question that will need to be addressed using data from multiple nuclear loci. Picture Information.

They-axis show the total percentage and the x-axis show the time in years. I genomes were successfully used to resolve long-standing phylogenetic questions, including the ratite and Elephantidae phylogenies.