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1980 Adesso Sono Te, Caduta Dellelt's critical
to see the Arizona Capitol dome from 1,500
feet in the sky. The GPS coordinates are Gila
Viewpoint, an overlook between Sedona and

Red Rock Crossing that's open to the public. No permit, no fee, and no waiting. Plenty of trail access, too, which is better than the alternative (fewer amenities, more concern for your safety). In *The Life of Arizona: Safaris, Treks, and Wilderness Adventures* (American City Guides, \$25), author Ross Brooks tells it like it is, in these 100 photos from Arizona's Grand Canyon and Four Corners region. This article appeared in the *Arizona Highways* May-June 2015 issue Click on the map to see other locations of this gallery. See more at: Photos

of Arizona Highway by: Ron Gieg |

Geowebphotos.com Pan-Indian gene tree method of human phylogeography The Pan-Indian gene tree method of human phylogeography (PI-MEG) is a method that attempts to reconstruct human population history by using the information derived from combining all existing data on uniparental genetic markers. The approach does not require either a large number of loci to be used in a concatenated alignment, nor a minimum of independent samples. The method provides a novel and theoretically

sound way to evaluate the validity of the clustering of different human populations as derived from mitochondrial, Y and autosomal markers. The method was proposed by the researchers F. Lage, R.A. Willing, R.T.W. Francis and H.C. Bunck in the paper Clustering versus Population Structure in Human Genetics, 2001. The paper was published in the Journal of Human Genetics. Clustering versus Population Structure in Human Genetics Population studies are often performed using data sets composed of samples taken from geographically isolated, linguistically homogeneous or genetically unrelated populations. The analysis of such population samples usually leads to "hard" clustering of individuals into distinct population groups, which may only be partially coincident with language families and geographic regions. The existence of such hard clustering is widely accepted, but the origins of the subdivision 6d1f23a050

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