Title of project:	Ancient mitochondrial DNA sequence and SNP data from Andaman and Nicobar museum samples
Name of database/ spreadsheet file:	Andaman_Results

Repeat the following section for each table within your database:

Name of table /	AndamanM31		
worksheet 1:			
Purpose of	Sequence data and SNP genotyping of Andaman M31 ancient samples		
table/worksheet:			
Number of rows	22		
of data:			
Primary key			
(database only):			
Name of field	Full description of field and sodes on towning lawy yeard	Data tuma and	
Name of field	Full description of field and codes or terminology used	Data type and field length (database only)	
Sample UIN	Unique Idenfifying Number for DNA extract		
Source	Source of Sample (NHM=Natural History Museum London,		
	OUMNH=Oxford University Museum of Natural History,		
	Duckworth = Duckworth Collection Cambridge)		
Population	Name of population		
Control Region	Results from cloned sequence data produced by PCR of		
Diversity	mtDNA control region		
Nucleotide	Results from SnapShot genotyping of mtDNA coding region		
Positions tested	sites tested (nt = not tested, nd = not determined)		
in Multiplex			
in Multiplex			

Name of table / worksheet 2:	AndamanM32		
Purpose of table/worksheet:	Sequence data and SNP genotyping of Andaman M32 ancient samples		
Number of rows of data:	34		
Primary key (database only):			
Name of field	Full description of field and codes or terminology used	Data type and field length (database only)	
Sample UIN	Unique Idenfifying Number for DNA extract		
Source	Source of Sample (NHM=Natural History Museum London, OUMNH=Oxford University Museum of Natural History, Duckworth = Duckworth Collection Cambridge)		
Population	Name of population		
Control Region Diversity	Results from cloned sequence data produced by PCR of mtDNA control region		
Nucleotide Positions tested	Results from SnapShot genotyping of mtDNA coding region sites tested (nt = not tested, nd = not determined)		

in Multiplex	