

Title of project:	Ancient mitochondrial DNA sequence and SNP data from Andaman and Nicobar museum samples
Name of database/ spreadsheet file:	Andaman_Non_M31_M32

Repeat the following section for each table within your database:

Name of table / worksheet 1:	A2_plex	
Purpose of table/worksheet:	Reporting results of A2 multiplex from Non_M32_M32 Andaman samples	
Number of rows of data:	6	
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	<i>Unique Identifying Number for DNA extract</i>	
<i>Source</i>	<i>Source of Sample (NHM=Natural History Museum London, Duckworth=Duckworth Collection Cambridge)</i>	
<i>Population</i>	Name of population	
Nucleotide Posiitons tested in A2 Multiplex	Results from SnapShot genotyping of mtDNA coding region sites tested (nt = not tested, nd = not determined)	

Name of table / worksheet 2:	B2_plex	
Purpose of table/worksheet:	Reporting results of B2 multiplex from Non_M32_M32 Andaman samples	
Number of rows of data:	6	
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	<i>Unique Identifying Number for DNA extract</i>	
<i>Source</i>	<i>Source of Sample (NHM=Natural History Museum London, Duckworth=Duckworth Collection Cambridge)</i>	
<i>Population</i>	Name of population	
Nucleotide Posiitons tested in A2 Multiplex	Results from SnapShot genotyping of mtDNA coding region sites tested (nt = not tested, nd = not determined)	