

Agharkar Research Institute, Pune, INDIA
MACS Collection of Microorganisms (MCM), WDCM 561

A Distinguished Repository of Anaerobes & Extremophiles

Form for Phylogenetic Analysis

1. User's Information			
a. Name & Designation of User			
b. Department / Division			
c. Name of Organization			
d. Is it Academic or Non-academic?			
e. Official Postal Address (with PIN Code)			
f. Email Address for Communication			
g. Contact Phone Number			
2. Details of Data Sent			
Sr. No.	Strain No./ Sequence Accession No.	Taxonomic Designation*	SRN (For MCM Use Only)
1.			
2.			
3.			
4.			
5.			
* Each phylogenetic tree (ARI Service No. 2.33) with up to 20 sequences will be considered a separate service. User may choose to combine strains/sequences with same genus designation into a single tree.			
3. Parameters for Phylogenetic Analysis			
Method:	Neighbor Joining	Maximum Parsimony	Do Not Know
Bootstrapping:	500 Iterations	1000 Iterations	Do Not Know
4. Payment Information			
a. Date & Mode of Payment			
b. UTR/Transaction ID for Payment			
5. Billing Information			
c. Name & Designation			
d. Address (with PIN Code)			

Organization Seal

Date & Signature of the User/ Authorized Signatory