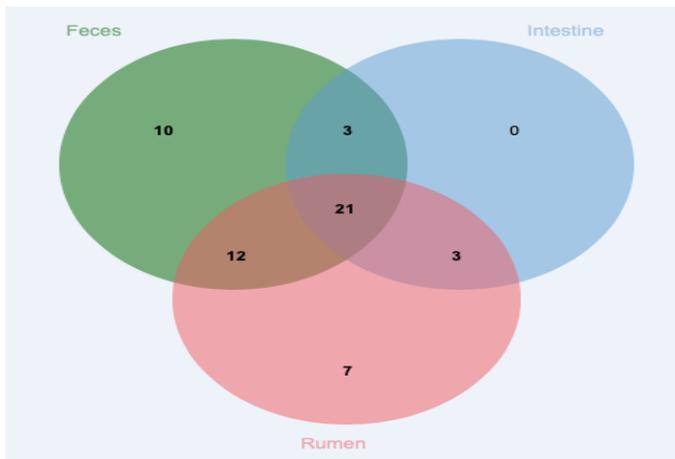


Supplementary Figure S1_Family core. Core microbiota at Family level of rumen, large intestine and feces from differentially fed water buffaloes. Families shared by at least 80% of samples included in each body site: (A) traditionally fed water buffaloes; (B) alternatively fed water buffaloes.

A Families included in the core microbiota in traditionally fed water buffaloes.



B Families included in the core microbiota in alternatively fed water buffaloes.

