



SDI Review Form 1.6

Editors comment		Author's comment (if agreed with reviewer, correct the manuscript and highlight that part in the manuscript. It is mandatory that authors should write his/her feedback here)
Compulsory REVISION comments	In the current study, the authors use regions in the mitochondrial genome (mtDNA) for determination of new sub-variations in the phylogeny of Egyptian sheep breeds. For this goal, modern nowadays' methods for assay are used, which gives a possibility for a maximally correct determination of the respective sub-variation belonging to the respective taxonomy unit. Besides for determination of phylogeny in sheeps, the ideas and methods, developed and described in the current research paper, could be applied for the same goals in every other species, including single-cell eukaryotic organisms (yeasts), single-cell prokaryotic organisms (bacteria strains), but also DNA- and RNA-viral strains, by small appropriate technique modifications for adaptation to the respective needs, which could be usable in a long-term period for development of methods about determination of appropriate places for transplantation, depending of the determined characteristics of cells, incubated in respective appropriate conditions for these goals	
Minor REVISION comments	The current research paper is a very important scientific contribution in the taxonomy point of view	
Optional/General comments	Sincerely recommend the current original research paper be published in the current issue of the scientific journal "Annual Research & Review in Biology" as it is	

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