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Poster Session 2 – Taxonomy Genetics

81 Maternal divergences within *Myospalax* and introgressive hybridization in the eastern Qinghai-Tibet Plateau

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The zokor (genus Myospalax) is a subterranean solitary species. It remains unknown what had caused the morphological complexity and difficulty in species circumscription within Myospalax. Hybridization and gene flow are proposed for explaining such a scenario, especially for ambiguous hybridization origin of Myospalax smithii. The morphological characters of this species suggested its relationship to both old and young species; however, hybridization usually occurs between young species or lineages with recent divergences. Our phylogenetic analyses of mitochondrial cytochrome b gene (Cyt b) and 12S rRNA gene obtained from 102 individuals representing all eight species of the genus Myospalax produced well discerning maternal phylogeny within the genus. Our results further rejected a hybridization origin hypothesis of Myospalax smithii between Myospalax cansus and Myospalax myospalaxi. However, we found that morphologically defined Myospalax smithii individuals interweaved with those belonging to Myospalax bailevi. These individuals from two species clustered into two well supported clades. We suggest that Myospalax smithii should be recognized as separate species, but the extensive hybridization and gene flow between it and Myospalax baileyi might have swamped out most pure Myospalax smithii individuals possibly due to heterosis of hybrids. Our results are consistent with the previous assumption regarding occurrence of hybridization between recently divergent lineages with close relationships. However, this conclusion was drawn based only on a combination of morphological and maternal evidence; further evidence with genetic signatures from both parents (e.g. nuclear genes) is needed. Overall, these results suggest that Myospalax provides a model system for studying speciation, reproduction isolation and gene flow of small mammals.

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