明治大学先端数理科学インスティテュート

MIMS顕像数理介了正位写十一

日時:2021年11月22日(月)(13:30 - 14:10)

場所:今年度はZoomでのリモート開催となります

Recent advancements of phylogenetic method.

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Abstract: Among biologists, the Phylogenetic Comparative Method (PCM) is an indispensable statistical tool to infer phenotypic evolution in a macro time scale (e.g. >1 million years). Nevertheless, its mathematical model has been quite simple because we cannot observe a past state of an organism, and thus strong assumptions are needed to yield a solvable problem. In this talk, I review recent methodologies of PCM and introduce a more flexible yet reliable approach based on a population genetics theory.



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