Dear Colleague,

We are pleased to invite you to attend the International Symposium “MODERN ACHIEVEMENTS IN POPULATION, EVOLUTIONARY AND ECOLOGICAL GENETICS (MAPEEG-2019)”. Meeting will be held by Far East Branch of the Russian Academy of Sciences (FEB RAS), National Scientific Center of Marine Biology FEB RAS (NSCMB, Vladivostok), Institute of Biology and Soil Science FEB RAS, Vladivostok City Administration, Far Eastern Federal University, and Vladivostok Public Foundation for Development of Genetics (VFDG).

The symposium MAPEEG-2019 is the tenth symposium held in Vladivostok and Vostok Marine Biological Station. The symposium will be arranged on September 8-13, 2019. Current meeting will be basically hosted by NSCMB.

The main subjects are the following:
- Evolutionary Genetics & Genomics
- Molecular Systematics, DNA Barcoding and Phylogenetics
- Microevolution. Population-Genetic Structure of Species
- Ecological Genetics

Some excursions and informal activities will be also developed during and after the meeting.

Organizing committee:
Yuri Kartavtsev (Chairman), Dmitry Atopkin, Sergey Turanov (Moderators), Anton Chichvarkin, Olga Chichvarkhina, Irina Kartavtseva, Oleg Katugin, Natalia Masalkova, Alexander Redin, Sergey Shedko, Irina Sheremetyeva, Anna Zolotova, Olesia Rutenko (Members).

For VFDG members no registration fee is offered; applying students may have some discount on the request. Working language at the meeting is English.

Registration forms will be accepted by March 20, 2019; abstracts by May 20, 2019 (1 page, Times NR 12, 1.0 space interval (the speaker’s name must underlined in the Registration Form). Samples of the Registration Form and Abstract are attached along with this circular.

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Sample of an abstract

TWENTY-TWO KEY ASPECTS TO THE PHYLOGEOGRAPHIC REVOLUTION IN POPULATION GENETICS

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Following its inception in the late 1970s and 1980s, the field of phylogeography has gradually transformed scientific thought and empirical approaches in population genetics and evolutionary biology. No longer must intraspecific evolution be viewed merely as a process of shifting allele frequencies within and among populations; population genetics can now be seen as a genealogical process also, extended through time and expanded across space. No longer must speciation be the line of demarcation below which phylogenetic perspectives have no relevance; instead, intraspecific evolution can now be interpreted as a genealogical coalescent process, and an organismal phylogeny can be seen as composed of multitudinous gene genealogies to which phylogenetic terminologies and concepts (properly adapted) can be applied. No longer must equilibrium scenarios (e.g., between mutation and selection, or between selection and drift) hold primary sway, simply for reasons of mathematical tractability, on population-genetic models; instead, the species-specific historical idiosyncrasies as well as generalities of real-life evolution can now be addressed explicitly. Finally, no longer need microevolutionary biology (including population genetics and ecology) be mostly divorced from macroevolutionary biology (including phylogenetics and systematics); the field of phylogeography now provides a powerful empirical and conceptual bridge between these traditionally separate evolutionary arenas. In this talk, I will justify and review more than 20 major aspects to what can rightly be termed the 'phylogeographic revolution' in population genetics.