

Press Release



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How eelgrass spread around the world

An international research team led by GEOMAR reconstructs the worldwide colonisation history of the most widespread marine plant

20.07.2023/Kiel/Groningen. Seagrasses are the only fully submerged, marine flowering plants to have conquered coastal habitats around the world. Eelgrass has captured the northern hemisphere, spanning latitudes from 35° to 70° North – from warm-temperate shores to frozen Arctic coasts. An international group of researchers coordinated by Professor Thorsten Reusch, marine biologist at GEOMAR Helmholtz Centre for Ocean Research Kiel, reconstructed the colonisation history of the eelgrass *Zostera marina* from its origin in the Northwest Pacific Ocean to the Pacific, Atlantic and the Mediterranean. In addition, they found a reduction in genetic diversity, which raises concerns how well eelgrass is able to adapt to a changing climate.

Seagrasses evolved from freshwater plants and use sunlight and carbon dioxide (CO₂) for photosynthesis and are able to thrive in depths down to 50 metres. In contrast to algae, they possess roots and rhizomes that grow in sandy to muddy sediments. The grass-like, leaf-shoots produce flowers and complete their life cycle entirely underwater. Seeds are negatively buoyant but seed-bearing shoots can raft, thus greatly enhancing dispersal distances at oceanic scale.

As a foundational species, eelgrass provides critical shallow-water habitats for diverse biotas and also provides numerous ecosystem services including carbon uptake. Seagrasses have recently been recognised as one of the important nature-based contributions to store carbon in the ocean. The sediment below seagrass meadows can sequester between 30 and 50 times more carbon annually than the roots of forests on land. Unfortunately, the continuing loss of seagrass beds worldwide including eelgrass is of acute concern.

An international group of researchers coordinated by Professor Thorsten Reusch, Head of the Research Division Marine Ecology at GEOMAR Helmholtz Centre for Ocean Research Kiel, used complete nuclear and chloroplast genomes from 200 individuals and 16 locations to reconstruct and date the colonisation history of the eelgrass *Zostera marina* from its origin in the Northwest Pacific Ocean to the Pacific, Atlantic and the Mediterranean. The findings described in a peer-reviewed publication and a Research Briefing published 4 days before the embargo end date (20 July 2023) are available here [W*](#)

assume that there were no eelgrass-based ecosystems hotspots of biodiversity and carbon storage in the Atlantic before that time. Recency was also mirrored in an analysis of the associated faunal community, which features many fewer specialised animals in the Atlantic as compared to the Pacific eelgrass meadows. This suggests that there was less time for animal-plant co-evolution, said Reusch. Mediterranean populations were founded from the Atlantic about 44 thousand years

ago, with western and eastern Atlantic shores only (re)expanded from refugia after the Last Glacial Maximum, about 19 thousand years ago and mainly from the American east coast with help from the Gulf Stream.

In addition, the researchers further confirmed the huge difference in genomic diversity between the Pacific and Atlantic, including latitudinal gradients of reduced genetic diversity in northern populations.

Atlantic eelgrass is less diverse on a genetic level than their ancestors by a factor of 35 among the most and least diverse populations, said Lei Yu, first author of the publication which was published in *Nature Plants*.

The study raises concerns as to how well Atlantic eelgrass, will be able to adapt to climate change and other environmental changes.

The loss of seagrass meadows at the southern range limits, in particular North Carolina and southern Portugal. In addition, heat waves have also caused losses in shallow waters.

because seagrass meadows form diverse and productive ecosystems, and no other species is able to take on the role of eelgrass if meadows cannot persist under future conditions.

The study fortifies diversity in the Atlantic. Our next step is to interrogate the eelgrass pangenome. A new reference genome from Pacific eelgrass is currently under development and should tell us more about the genetic diversity of eelgrass.

The study was led by Dr. Reusch, emeritus professor from the University of Groningen who initiated the study and coordinated the work between the Joint Genome Institute (JGI) and the research team. Thus, the verdict on rapid genetic divergence is clear.

Original publications:

Yu, L. et al (2023): Ocean current patterns drive the worldwide colonization of eelgrass (*Zostera marina*). *Nature Plants*, doi: [10.1038/s41477-023-01464-3](https://doi.org/10.1038/s41477-023-01464-3). The publication will become available on 15 November 2023.