

Northeast Microbial Database: a web-based databank of culturable soil microbes from North East India

S. R. Joshi*, Subhro Banerjee, Kaushik Bhattacharjee, Nathaniel A. Lyngwi, Khedarani Koijam, Polashree Khaund, Lamabam Sophiya Devi and Fenella M. W. Nongkhlaw

Microbiology Laboratory, Department of Biotechnology and Bioinformatics, North-Eastern Hill University, Shillong 793 022, India

Northeast Microbial Database (NEMiD) is a first of its kind digital database on microbial diversity from North East (NE) India, which provides information on culturable microbes (bacteria, fungi and actinomycetes) isolated from soils of the region (Arunachal Pradesh, Assam, Manipur, Meghalaya, Mizoram, Nagaland, Tripura, Sikkim and parts of North Bengal) distributed across landscape elements (LSE) spread in various ecosystems. The database is based on the primary data through survey, isolation and characterization of the microbes during 2008–2012. Details about a particular microorganism in the database cover aspects of phylogenetic information on taxonomic outline and classification, morphological, biochemical and molecular characterization, general and cultural characteristics, geographical/location information, history of cultures and preservation details. Each microbe is identified based on cultural, biochemical and molecular characteristics leading to its taxonomic position up to the species level along with information on sampling sites comprising altitude, soil type, latitude, longitude, soil temperature, ambient temperature, humidity and soil pH. The database is available on-line at <http://mblabnehu.info/nemid/>.

Keywords: Digital database, landscape elements, natural ecosystems, soil microbes.

MICROORGANISMS play a vital role in the maintenance of the earth's ecosystems. Microbes have mainly remained unused due to poor documentation. At the global level, databases on microbes that have been developed include Species 2000 and the Integrated Taxonomic Information System (ITIS) collaborative Catalogue of Life (www.gbif.org)¹. Many Microbe Microarrays Database (M³D) is a resource for analysing and retrieving gene expression data for microbes. MicrobeID.com is an on-line resource for microbiologists, where one can find information related to identification keys, databases, test methods (including images and video), media guides and other useful resources on microbes (<http://www.microbeid.com>).

A database providing comprehensive information about 600 known bacteria living in human mouth can be found at <http://homd.org/>.

Few fungal databases are available on the World Wide Web and they include; Systematic Botany and Mycology Laboratory (SBML), US Department of Agriculture, Agricultural Research Service, Betsville, Maryland USA (<http://nt.ars-grin.gov>); MICH Fungal Bioinformatics: A Myclopedia Project at University of Michigan Herbarium, USA (<http://www.herb.lsa.umich.edu/bioinformatics.html>); Catalog of Rust Types, New York Botanical Garden (<http://www.nybg.org/bsci/hcol/rust/pucci1.html>); The Pen State Mycological Herbarium, USA (<http://www.ma.psu.edu/pacma>); and the microbial database of Virginia Bioinformatics Institute (VBI) which hosts data from a range of plant pathogenic oomycetes, fungi and bacteria primarily those under the VBI (<http://www.vbi.vt.edu>).

In all the above efforts, documentation has been attributed to research in the US. In India, some initiatives have been made, but overall data on Indian microbial diversity are negligible and no microbial documentation has so far been done for North East India. There is paucity of information that links microbes to various ecological systems and their potential uses in our country. In India, the Environmental Information System (ENVIS) Centre, Department of Zoology, University of Madras has developed databases on thematic areas such as reference/abstract database, experts' directory database of microbiologists, bibliographic database and database on research activities on microbes (Ph D thesis, research projects, reports, monographs/manuals). Further, the centre has created a database on the biodiversity of microbes (isolation, habitat, ecological significance), especially marine microbes (<http://www.envismadrasuniv.org/index.php>). The Department of Biotechnology in Presidency College, Chennai has developed a database on the bioinformatics of antibiotics from microbes and another on vaccines is being developed (<http://www.bifcpresidency.tn.gov.in/database.html>). The Alagappa University, Karaikudi, Tamil Nadu generated a database for the 16S rRNA sequences of marine bacteria from the Gulf of Mannar for biotechnological prospects (<http://www.alagappabiotech.org/biobank/biobank%20index%20page.htm>). The Bioinformatics Center of the Institute of Bioresources and Sustainable Development, Imphal is developing a biodiversity database on cyanobacteria of NE India².

The present web-based NEMiD database is based on data generated for culturable microbes (Figure 1) from surface and subsurface soils collected across different altitudinal zonations of NE India and parts of North Bengal. The sampling regions were located between 20°N and 32°N lat. and 87°E and 100°E long. and occupy an area of 262,230 sq. km. The database accessible at <http://mblabnehu.info/nemid/>. Soil samples were collected from different microhabitats, mixed thoroughly to make a composite sample for microbial analysis (Table 1).

*For correspondence. (e-mail: srjosshi2006@yahoo.co.in)

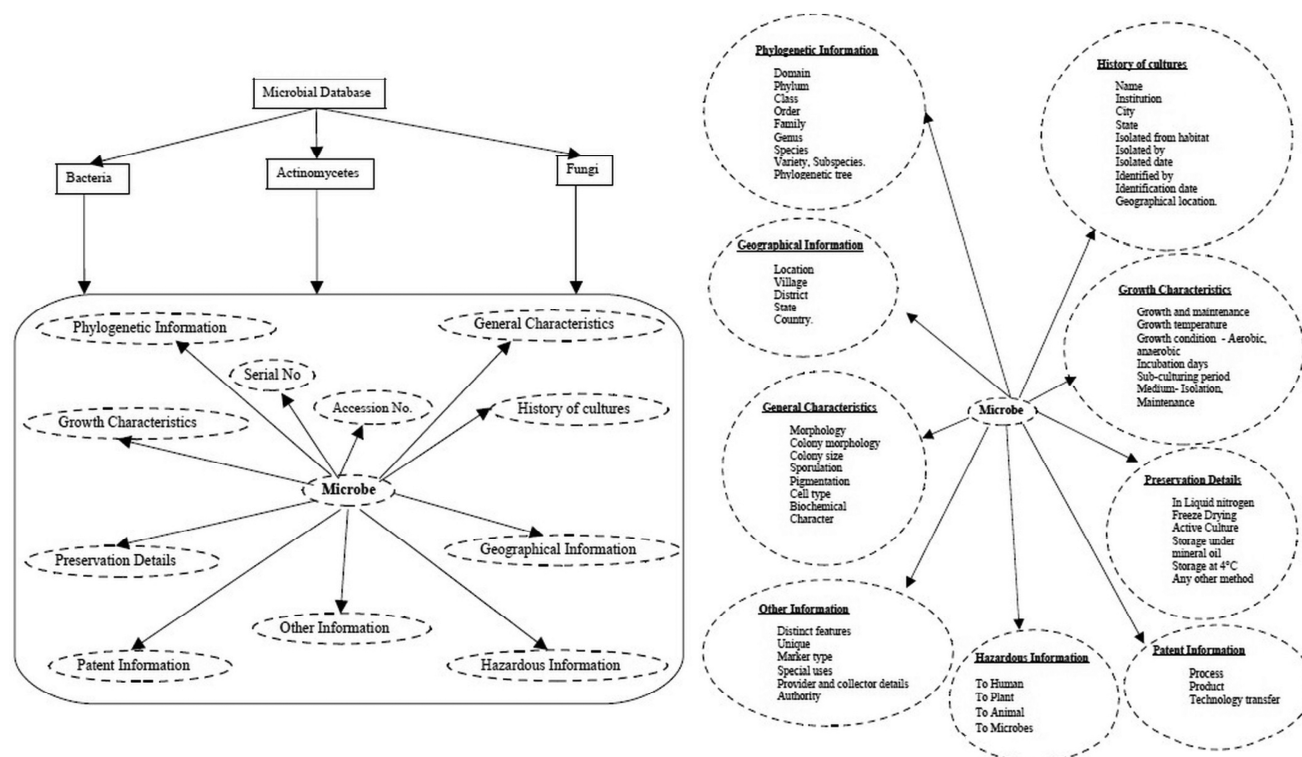


Figure 1. Interface of database parameters.

Table 1. Parameters used for identifying the microbes incorporated in the database

Identification characteristics	Parameters/tests
Morphological	Fungi: Colony morphology, hyphae, asexual spores, sexual spores, reproductive bodies and arrangement of conidia. Bacteria: Cultural characteristics (colour, form, margin, elevation, appearance), Gram-staining. Actinomycetes: Colony characteristics (colour, margin, elevation, presence of substrate/aerial mycelium), staining (simple stain, Gram-stain, acid alcohol fast stain), pleomorphism, endospore detection and motility.
Biochemical	Bacteria: IMViC, catalase, oxidase, amylase, lipase, gelatinase, nitrate reduction, triple sugar iron test, urease, casein hydrolysis, cellulase, protease, H ₂ S production, N ₂ gas production, coagulase test, sugar utilization (mannitol, arabinose, dextrose, sucrose, xylose, rhamnose, lactose, maltose), arginine dihydrolase, phenylalanine deaminase, amino acid decarboxylase (lysine, ornithine), O/F test. Actinomycetes: Allantoinase, nitrate reduction, nitrite reduction, phosphatase, oxidase, urease, indole, Voges-Proskauer, methyl red, catalase, starch hydrolysis, gelatin hydrolysis and casein utilization.
Molecular	Phylogenetic analysis based on 18S rRNA/5.8S rRNA/16S rRNA/ITS/IGS sequences.

Microbial identification is an intensified process to include all the NE states on uniform basis. Various parameters like latitude and longitude, altitude, ambient temperature, humidity, soil temperature, vegetation and soil types were noted for further references (Table 2). The data collection on microbes is an ongoing process carried out uniformly across all the states of NE India and the microbes are incorporated in the database as and when the new isolates are characterized from the sites. The diversity is expected to vary significantly with an increase in the number of samples, which is being documented and updated in the database.

Microorganisms inhabit virtually every niche in the biosphere and they are the most adaptable group of organisms that are able to survive almost all conditions. They form a vast community with a diversity of protozoa,

algae, fungi, bacteria and viruses that are associated with all plants and animals, also being responsible for their well-being³. Many of them are beneficial to human beings, and are used in health-care products, drug industry, food industry and agriculture⁴⁻⁸.

Soil is the natural habitat for all organisms involved in important ecological processes. Soils are excellent cultural media for the growth of many types of organisms, including bacteria, fungi, algae, protozoa and viruses. In addition, various nematodes and insects are also present in the soil⁹. A spoonful of soil contains billions of microorganisms. In general, majority of microbial population is found in the upper 12 inches of soil and their number decreases with depth¹⁰. Majority of bacteria, however, are heterotrophic and utilize large amounts of organic matter. The development of fungi is especially favoured by soils

RESEARCH COMMUNICATIONS

Table 2. Interface for landscape elements and data search in NEMiD

Landscape elements/physiographic factors	Latitude, longitude, altitude, relative humidity, ambient temperature, soil temperature, soil pH, soil depth, vegetation type, soil organic content.
Species distribution map	Geographical locations of microbial isolates on the map of North East India using Google Maps.
Isolate details	Classification of the isolates under the domain Actinomycetes, Bacteria, Fungi, which includes parameters like phylogenetic information covering the taxonomic outline and classification, morphological, biochemical and molecular characterization, general and cultural characteristics, geographical/location information, history of cultures, preservation details, hazardous information, patent information, detailed identification, phylogenetic information covering the taxonomic outline, morphological, biochemical and molecular characterization, general and cultural characteristics, geographical/location information, history of cultures, preservation details, hazardous information, patent information.
Score sheet details	State, district, sampling site name, sampling site code, soil temperature, ambient temperature, humidity, moisture content, soil pH, soil depth, soil type, soil texture, vegetation, longitude, latitude, altitude, collection time (IST), collection date, season, nearest village, landmark collector(s).



Figure 2. Snapshot of user interfaces available in the database.

having an acidic reaction and an aerobic condition near the surface. As fungi exist in both the mycelial and spore stages, it is difficult to enumerate them^{11,12}.

Microorganisms have opened vistas for biotechnological applications and have been explored for the discovery of new pharmaceutical products, synthesis of new enzymes, chemicals and new organisms that carry out novel processes¹³. A study of the biodiversity of microorganisms and their potential applications provides scope for development of new strains through genetically engineered microbes/recombinant DNA technology. These unique combinations may be able to degrade complicated pollutants prevalent in the environment¹⁴.

NEMiD is unique and is specific to the microbial diversity of NE India (Figure 2), which is likely to open avenues for its exploration for human well-being.

Further, NE India being one of the mega biodiversity hotspots of the world, has natural ecosystems still prevalent in pristine conditions offering scope for bioprospection of novel organisms hitherto unknown to the scientific world. Further, the sacred groves in this region are rich patches of undisturbed virgin forests and serve as a natural habitat for many endemic, rare and economically valuable species¹⁵. There have been efforts to document the fauna and flora of the region, but documentation of microbes from the region has been scarce. Therefore, a comprehensive database on microbes of NE India is complementary to the development of such a database at the country level in future. The present database incorporates data on microbes isolated from soil samples collected uniformly from the NE states of India (Figure 3).

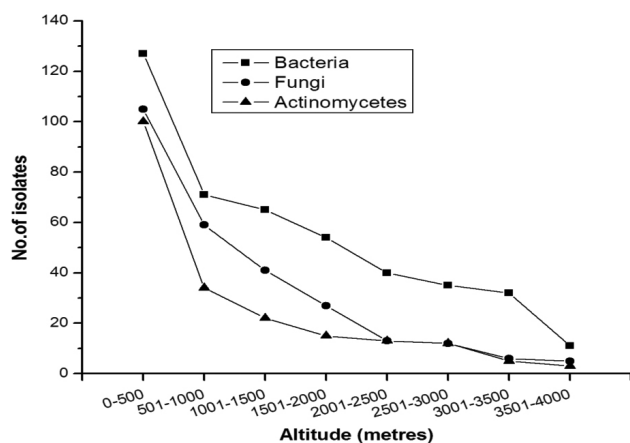


Figure 3. Variation in the number of isolates (bacteria, fungi, actinomycetes) with altitude.

As the enormity of microbial diversity becomes apparent, the dilemma of how to preserve microorganisms and their gene pools asserts itself ever more forcibly¹⁶. *Ex situ* preservation in culture collections can never be expected to hold more than a small fraction of the genetic diversity of microorganisms. Consequently, *in situ* conservation is as crucial for the protection of microbial diversity as it is for macrobiological diversity, as conserving biodiversity equals conserving ecosystems¹⁷. In recent years, the frequency of recommended strategies for the conservation of biodiversity has increased swiftly; few, however, pay attention to microorganisms. The characterized microbes are preserved by freezing at -80°C to maintain viability and reuse. Majority of the representative isolates have been deposited at Institute of Bioresources and Sustainable Development (IBSD), Imphal, India.

The occurrence of life forms in a region can be linked to many ecological factors. The factors and habitats in NE India are well known to be decisive in microbial diversity distribution^{18–35}. The present database holds data on soil microbes generated through primary data collection undertaken for various niches of NE India. This is an ongoing project and data on additional microbes are being documented and incorporated into the database as and when new isolates are characterized.

- Palacio, M., Towards an index of all known species: the catalogue of life, its rationale, design and use. *Integr. Zool.*, 2006, **1**, 18–26.
- Tiwari, O. N., Oinam, G., Koijam, L., Devi, S. D. and Singh, O. A., Cyanobacterial biodiversity, conservation and possible commercial exploitation of Indian region falling Indo-Burma biodiversity hotspot. *Biosci. Biotechnol. Res. Commun.*, 2009, **1**(2), 5–32.
- Sota, T., Interactions among microorganisms, animals and plants. 17th Symposium of the Society of Population Ecology. *Res. Popul. Ecol.*, 1996, **38**(2), 183–184.
- Leroy, F. and De Vuyst, L., Lactic acid bacteria as functional starter cultures for the food fermentation industry. *Trends Food Sci. Technol.*, 2004, **15**(2), 67–78.
- Demain, A. L., Small bugs, big business: the economic power of the microbe. *Biotechnol. Adv.*, 2000, **18**(6), 499–514.

- Leeds, J. A., Schmitt, E. K. and Krastel, P., Recent developments in antibacterial drug discovery: microbe-derived natural products – from collection to the clinic. *Expert. Opin. Invest. Drugs*, 2006, **15**(3), 211–226.
- Berg, G., Plant–microbe interactions promoting plant growth and health: perspectives for controlled use of microorganisms in agriculture. *J. Appl. Microbiol. Biotechnol.*, 2009, **84**(1), 11–18.
- Antoni, D., Zverlov, V. V. and Schwarz, W. H., Biofuels from microbes. *J. Appl. Microbiol. Biotechnol.*, 2007, **77**(1), 23–35.
- Fitter, A. H., Gilligan, A. A., Hollingworth, K., Kleczkowski, A., Twyman, R. M. and Pitchford, J. W., Biodiversity and ecosystem function in soil. *Funct. Ecol.*, 2005, **19**, 369–377.
- Blumea, E., Bischoff, M., Reichert, J. M., Moormann, T., Konopka, A. and Turco, R. F., Surface and subsurface microbial biomass, community structure and metabolic activity as a function of soil depth and season. *Appl. Soil Ecol.*, 2002, **20**(3), 171–181.
- Pepejnjak, S. and Segvic, M., Occurrence of fungi in air and on plants in vegetation of different climatic regions in Croatia. *Aerobiologia*, 2003, **19**(1), 11–19.
- Wu, Z., Tsumura, Y., Blomquist, G. and Wang, Xiao-Ru, 18S rRNA gene variation among common airborne fungi, and development of specific oligonucleotide probes for the detection of fungal isolates. *Appl. Environ. Microbiol.*, 2003, **69**(9), 5389–5397.
- Demain, A. L., Microbial biotechnology. *Trends Biotechnol.*, 2000, **18**(1), 26–31.
- Sayler, G. S. and Ripp, S., Field applications of genetically engineered microorganisms for bioremediation processes. *Curr. Opin. Biotechnol.*, 2000, **11**(3), 286–289.
- Samati, H. and Gogoi, R., Sacred groves in Meghalaya. *Curr. Sci.*, 2007, **93**(10), 1338–1339.
- Smit, E., Leeftang, P., Glandorf, B., Van Elsland, J. D. and Wernars, K., Analysis of fungal diversity in the wheat rhizosphere by sequencing of cloned PCR-amplified genes encoding 18S rRNA and temperature gradient gel electrophoresis. *Appl. Environ. Microbiol.*, 1999, **65**, 2614–2621.
- Lomolino, M. V., Elevation gradients of species density: historical and prospective views. *Global Ecol. Biogeogr.*, 2001, **10**, 3–13.
- Devi, L. S. and Joshi, S. R., Antimicrobial and synergistic effects of silver nanoparticles synthesized using soil fungi of high altitudes of eastern Himalaya. *Mycobiology*, 2012, **40**(1), 27–34.
- Bhattacharjee, K., Banerjee, S. and Joshi, S. R., Diversity of *Streptomyces* spp. in Eastern Himalayan region – computational RNomics approach to phylogeny. *Bioinformatics*, 2012, **8**(12), 548–554.
- Banerjee, S., Rai, S., Sarma, B. and Joshi, S. R., Bacterial biofilm in water bodies of Cherrapunjee: the rainiest place on planet earth. *Adv. Microbiol.*, 2012, **2**(4), 465–475.
- Saikia, P. and Joshi, S. R., Changes in microfungus community in Cherrapunji – the wettest patch on earth as influenced by heavy rain and soil degradation. *Adv. Microbiol.*, 2012, **2**, 456–464.
- Devi, L. S., Khaund, P., Nongkhaw, F. M. W. and Joshi, S. R., Diversity of culturable soil micro-fungi along altitudinal gradients of eastern Himalayas. *Mycobiology*, 2012, **40**(3), 151–158.
- Lyngwi, N. A., Koijam, K., Sharma, D. and Joshi, S. R., Culturable bacterial diversity along the altitudinal zonation and vegetation range of tropical Eastern Himalaya. *Rev. Biol. Trop.*, 2013, **61**(1), 467–490.
- Banerjee, S. and Joshi, S. R., Insights into cave architecture and the role of bacterial biofilm. *Proc. Natl. Acad. Sci., India, Sect. B*, 2012, **83**(3), 277–290.
- Sarma, B., Acharya, C. and Joshi, S. R., Plant growth promoting and metal bioadsorption activity of metal tolerant *Pseudomonas aeruginosa* isolate characterized from uranium ore deposit. *Proc. Natl. Acad. Sci., India, Sect. B*, 2013, **84**(1), 157–164.
- Saikia, P. and Joshi, S. R., A study on the occurrence of non-O157 Shiga toxin producing *Escherichia coli* isolates in retail chicken

- meats marketed in North-East India. *Proc. Natl. Acad. Sci., India, Sect. B*, 2013, **84**(2), 337–342.
27. Kumar, R., Nongkhlaw, M., Acharya, C. and Joshi, S. R., Growth media composition and heavy metal tolerance behaviour of bacteria characterized from the sub-surface soil of uranium rich ore bearing site of Domiasiat in Meghalaya. *Indian J. Biotechnol.*, 2013, **12**, 115–119.
 28. Bhattacharjee, K. and Joshi, S. R., Phylogenetic rearrangement of *Streptomyces* spp. on the basis of internal transcribed spacer (ITS) region using molecular morphometrics approach. *Indian J. Biotechnol.*, 2013, **12**, 67–79.
 29. Kumar, R., Nongkhlaw, M., Acharya, C. and Joshi, S. R., Bacterial community structure from the perspective of the uranium ore deposits of Domiasiat in India. *Proc. Natl. Acad. Sci., India, Sect. B*, 2013, **83**(4), 485–497.
 30. Devi, L. S., Bareh, D. A. and Joshi, S. R., Studies on biosynthesis of antimicrobial silver nanoparticles using endophytic fungi isolated from the ethno-medicinal plant *Gloriosa superba* L. *Proc. Natl. Acad. Sci., India, Sect. B*, 2013; DOI:10.1007/s40011-013-0185-7.
 31. Bhattacharjee, K., Banerjee, S., Bawitlung, L., Krishnappa, D. and Joshi, S. R., A study on parameter optimization for degradation of endosulfan by bacterial consortia isolated from contaminated soil. *Proc. Natl. Acad. Sci., India, Sect. B*, 2013; DOI: 10.1007/s40011-013-0223-5.
 32. Khaund, P. and Joshi, S. R., Wild edible macrofungal species consumed by the Khasi tribe of Meghalaya, India. *Indian J. Nat. Prod. Resour.*, 2013, **4**(2), 179–204.
 33. Fenella, M. W. N. and Joshi, S. R., Micrographical assessment of antifungal effect of endophytic bacteria. *Proc. Natl. Acad. Sci., India, Sect. B*, 2014; DOI:10.1007/s40011-014-0321-z.
 34. Fenella, M. W. N. and Joshi, S. R., Distribution pattern analysis of epiphytic bacteria on ethnomedicinal plant surfaces: a micrographical and molecular approach. *J. Microsc. Ultrastruct.*, 2014; DOI:10.1016/j.jmau.2014.02.003.
 35. Bhattacharjee, K. and Joshi, S. R., NEMiD: A web-based curated microbial diversity database with geo-based plotting. *PLoS ONE*, 9(4), e94088.

Received 27 May 2014; revised accepted 18 January 2015

The May 21st, 2014 Bay of Bengal earthquake: implications for intraplate stress regime

A. K. Rai^{1,*}, S. Tripathy¹ and S. C. Sahu²

¹School of Earth, Ocean and Climate Sciences, Indian Institute of Technology Bhubaneswar, Bhubaneswar 751 007, India

²India Meteorological Department, Bhubaneswar 751 020, India

The northeastern part of the Indian Ocean, i.e. the Bay of Bengal (BoB) is located near some of the most complicated tectonic zones on the Earth. An earthquake of magnitude ~ 6.0 occurred on 21 May 2014 near the coast of Odisha. Occasional moderate to large earthquakes in BoB highlight the need to study precise hypocentre locations, and focal mechanisms to understand the cause of intraplate seismicity in BoB.

*For correspondence. (e-mail: akr Rai@iitbbs.ac.in)

It is also important for seismic hazard and tsunami risk evaluation along the eastern coast of India. We present an analysis of the digital data of this earthquake recorded by regional and global networks of seismic stations. Our analysis of travel-times of *P*- and *S*-waves indicates that the epicentre of the earthquake is located between the Eighty Five East and Ninety East ridges. The focus of this earthquake was at a depth of ~ 61 km, well below the lower boundary of the oceanic crust. The focal mechanism determined by modelling long period *P*- and *SH*-waveforms suggest a strike-slip motion along a NW–SE or NEE–SWW-directed fault or fracture. We interpret that the upper part of the BoB lithosphere is abnormally strong and brittle.

Keywords: Earthquake location, focal mechanism, intraplate seismicity, tectonic zones.

THE Indian Ocean is one of the most seismically active oceanic regions where intraplate seismicity is observed frequently¹ (Figure 1). The northeastern part of Indian

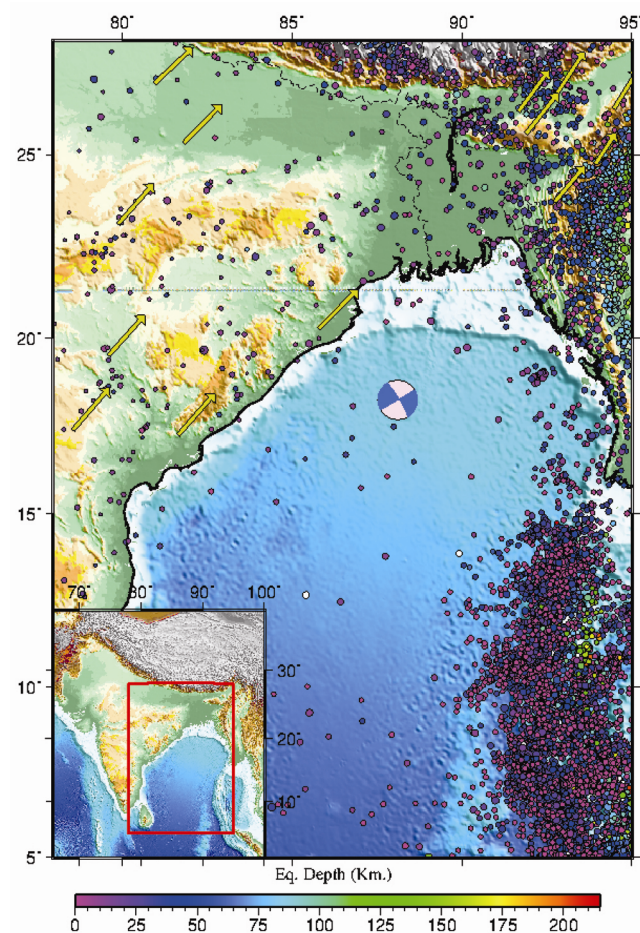


Figure 1. The location map and regional seismicity of the Bay of Bengal. The coloured circles represent earthquakes, where colour and size of a circles correspond to the depth, and magnitude of the earthquakes. Location of the 21 May 2014 earthquake is shown by beachball. Arrows show GPS velocity vectors³³.