

Climate change is real: fungal perspective!

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ABSTRACT

Climate change is apparent around the globe as it is dramatically affecting the natural ecosystems. The preliminary cause of climate change is increased green house gases, deforestation, and anthropogenic activities and the resultant change is seen in the form of extreme events like increased temperature, rainfall, permafrost thaw, glacial retreats, sea level, greater occurrence of wildfires, etc (NASA, 2016). It is causing tremendous effect on biodiversity therefore it becomes a serious concern globally. At the face of climate change the macroscopic organisms are either extinct or are at the verge of extinction and may adapt themselves for survival; efforts are successfully laid to conserve them but also it is equally important to focus on the existence, adaptation and alterations taking place in community composition of microscopic world. Impact of 1 trillion microbes inhabiting the planet earth on climate can be huge as they can accelerate climate change, exacerbated by pollution and habitat loss. However, they are understudied on climatic front, especially fungi, which represents a major portion of eukaryotic kingdom, resides in diverse habitats, constitutes considerable biomass on planet earth and regulate soil carbon feedback loop by deriving biogeochemical cycles; therefore they could be vulnerable to climate change largely by habitat loss. This review is an effort to draw the attention towards how fungal diversity responds to climate selection pressure either by adaptation or shifts in community composition observed during weather extremes.

KEYWORDS: Fungi, climate change, adaptation, green house gas, temperature

1. GLOBAL CLIMATE CHANGE

1.1. Climate change: Magnitude: Climate change is a rather slow and degrading process which is not homogeneous and its repercussions will be observed by mankind in the coming years. As Ram Ramanathan, the first person to predict global warming, rightly prefers the term 'climate disruption' over 'climate change' as it accurately points to the severity associated with it (Madden and Ramanathan, 1980). National agencies like NASA (National aeronautics and space administration) and NOAA (National oceanic and atmospheric administration) clearly forecast the magnitude at which climate is changing indicating ocean acidification that is increasing 2 billion tons per year, ocean warming which is more than 0.4 °F since 1969 to record levels, ocean acidification, glacial retreat, shrinking ice sheets, frequent wildfires and declining sea ice (<https://climate.nasa.gov/evidence/>). The global surface temperature in the year 2018 was the highest and was the fourth consecutive warmest year since 1880 and there is no wonder now the year 2019 embarks on a new history. Observing this rapid change of desertification around the world, Sudan is now declared uninhabitable and as one of the most vulnerable countries in the world to climate change (www.climatelinks.org).

Climate change is partly blamed in frequent devastations like wildfires of Colorado, drought occurring in Sudan and hurricane Maria hitting Puerto Rico in 2017 which makes the subject 'climate change' a pressing concern. What are the effects of climate change? Is it just the rise in temperature, water table, atmospheric carbon dioxide and ozone depletion causing a fast decline of macroscopic world or is it also adversely affecting the unseen majority of microscopic world. Climate change confers its deleterious effects equally to all five kingdoms of life, however, the microbes draw lesser attention at the face of climate change and are understudied in the context of conservation strategies. There have been lesser known facts available about their consequence on the belowground diversity and a connection is poorly established

as to how do they retaliate towards nature. In this review we will discuss this undermined area of belowground diversity in climate changed scenario with a special focus on fungi and how these microscopic organisms retaliate towards climate change in the macroscopic world.

1.2. Climate change: Prediction: According to IPCC (International Panel on Climate Change) 2013, global warming effects are heterogeneous like alpine, boreal and arctic areas are expected to experience four times greater warming and more variable precipitation than equatorial and temperate regions predicted to increase by 2 °C whereas EPA (Environmental Protection Agency), 2016 predicts the global mean temperature to increase by at least 2.7 °F within the next century. Carbon dioxide level tends to surpass 550 ppm in the next 30-80 years which will have detrimental effect on the nutritional value of food crops (Smith and Myers, 2018), likely range of global temperature increase is 2.04-9 °C, with median 3.2 °C and a 5 % chance that it will be less than 2 °C (1.5 °C) by 2100 (Raftery *et al.*, 2017).

1.3. Climate change: a serious concern: Worldwide protests against climate change are taking place and a group like Extinction rebellion declare them as 'unprecedented global emergency' that the humanity is facing at present; highlighting the importance of subject to policy makers (<https://rebellion.earth/wp-content/uploads/2019/04/REBEL-STARTER-PACK-14-March-2019.pdf>). Climate change becomes a serious concern when its impact is observed in all kingdoms of life and anthropogenic activities are largely responsible for it. According to IPCC anthropogenic activities have a measurable effect on the climate change for instance, global tourism alone accounts for about 8% of global green house emission (Lenzen *et al.*, 2018). However, land ecosystem absorbs 30% of human carbon dioxide emissions otherwise climate change would have been more rapid than it is thought to be. Recently in Australia, *Bramble Cay melomys* has been officially declared extinct by IUCN in February, 2019 and the reason behind the

first documented extinction of a mammal species involves climate change due to 'anthropogenic activities' (Woinarski and Burbidge, 2016). Such events are also intensifying the rapid outbreaks of fungal disease and their dispersal and thus threatening the earth's biodiversity. A striking example of detrimental rise in chytrid fungus *Batrachochytrium dendrobatidis* has led to an extinction of more than 200 sp. of frogs; their transmission is largely facilitated by humans (Scheele *et al.*, 2019) and rapid spread of *Candida auris* in more than 30 countries is alarming (Casadevall *et al.*, 2019). The devastating effect is often measured by the lives lost, effect on marine life includes ocean acidification, effect on wildlife, extinction risk associated with climate change but surprisingly effect on microbial world is seldom discussed.

Are only macrobes at risk of extinction? No, microscopic organisms too are vulnerable to climatic changes that may lead them to extinction, therefore worthy of conservation. Microbes are always assumed to be everywhere but few are particularly found only in specific habitats. For e.g. an anaerobic fungus inhabits in hindgut of the critically endangered Somali wild ass and is now at extinction risk through habitat loss even before it has been formally described. In addition, their interdependence on each other for survival may cause extinction risk to host species (Liggenstoffer *et al.* 2010). Microbes are often not a separate entity but are an integral part of food webs so that the extinction of one species puts the next species in taxon inadvertently to higher risk. In 2019, Hawaii forests faced the extinction of a rare species of tree snail (*Achatinella apexfulva*) which is linked to phyllosphere fungi. While feeding on fallen leaves they reduce fungal abundance which protect the tree (*Ohi'a*) from fungal diseases caused by *Ceratocystis*. Extinction of fungi will not only reduce a member from their kingdom but also affect other species linked to it. Many organisms are dependent on fungi for their nutrition and are species-specific in selecting them as their source of nutrition for e.g., *Collembula* population is strongly determined by elevated temperature when grazing on fungal species, *Resinicium bicolor* and *Phallus impudicus* (A'Bear *et al.*, 2012). Such findings should be given critical importance and are worthy of providing proper conservation strategies, attention and protection since fungi are dominant decomposers and beneficial in several other ways for the ecosystem.

2. FUNGAL DIVERSITY

Fungi in nature exist as major driver of key biogeochemical cycles involved in the process of decomposition and thus contribute 1.7×10^{27} Mb of DNA (Landenmark *et al.*, 2015) and 12 Gt of carbon biomass is reported to be encompassed by fungi on the planet earth (Bar-On *et al.*, 2018). For example fungi significantly contribute towards about 50-70% of carbon stored in boreal forest of Scandinavia so that this major portion accounts for the emission and regulation of carbon dioxide from soils (Clemmensen *et al.*, 2013). Moreover, shift in the fungal community structure due to climate change that may favour fungi over other microbes shall have dramatic impact on emission or sequestration of carbon dioxide and accelerating the climate change (Treseder

et al., 2016). Microcosm experiment reveals that AMF (Arbuscular Mycorrhizal Fungi) can affect soil and plant stoichiometry under global change of warming and nitrogen pollution; can reduce phosphorus limitation caused by nitrogen input and slow down negative influence of global change on plant growth (Mei *et al.*, 2019).

Fungi belong to the largest eukaryotic kingdom encompassing millions of species containing approximately 1,44,000 species which are formally named to date (https://stateoftheworldsfungi.org/2018/reports/SOTWFungi_2018_Full_Report.pdf). As is evident from various studies, fungi are inevitably present everywhere and attempts to enumerate their numbers and taxonomic diversity abound in literature (Bajpai *et al.*, 2019). An early attempt by Hawksworth (1991) estimated fungal diversity conservatively at 1.5 million species; subsequently 1.5 million was considered as a working hypothesis (Hawksworth, 2001). Further studies have suggested an altered ratio estimating fungal forms to 3.5 to 5.1 million species (O'Brien, 2005; Blackwell, 2011). Some of these uncertainties are a result of hidden fungal forms in plant world which is largely influenced by geographical location and edaphic factors. Tedersoo *et al.* (2014) conducted a study of 365 sampling sites globally and stated that the fungal diversity is overestimated by 1.5 to 2.5 times. Therefore, this varying ratio has led Hawksworth and Lücking (2017) to revisit the actual diversity of fungi to be within the range of 2.2 to 3.8 million species. Key drivers involved in controlling the fungal diversity act either directly or indirectly to exert their influence. Direct factors include environmental factors such as temperature and soil moisture. However, former has a positive effect at global and local scale whereas latter has a positive effect at local scale (Allison and Treseder, 2011). Indirect influences include aboveground community of plants which are linked through mycorrhizal interaction, utilize root exudates and are chiefly involved in decomposition process of plant litter. Therefore plant in many ways determines the belowground fungal communities.

3. SHIFTS IN FUNGAL COMMUNITY

Here we exemplify climate change, taking examples of few major ecosystems that are rich in microbes and are considered to be the largest in terms of sequestering carbon from atmosphere. As change in the below ground diversity may impact the future climate change therefore, shifts in such communities of micro-organism become important parameter to predict the climate change.

3.1. Soil ecosystem: Nearly 90% of plants on terrestrial ecosystem are associated with mycorrhizal fungi that helps them acquire nitrogen from soil and thereby improves plant performance. Ectomycorrhiza is a group of fungi that helps plant grow efficiently under high carbon dioxide levels (Terrer *et al.*, 2016) often termed as 'carbon dioxide fertilization effect'. Plants need nitrogen to respond to high levels of carbon dioxide in the atmosphere and therefore they need a readily available source of nitrogen. Weber *et al.* (2011) studied six different terrestrial ecosystem subjected to elevated carbon dioxide concentration having around 70% unique species to a given site. High fungal abundance was

observed in Creosote, Aspen and Marsh whereas Loblolly and scrub supported low fungal abundance. Abundance of *Polysporales*, *Basidiomycota*, *Pleosporales* and *Ascomycota* in carbon dioxide elevated conditions was obvious whereas *Amanita manicata* and *Chaetomium thermophilum* decreased in abundance.

Soil ecosystems are frequently flooded with nitrogenous fertilizer which may inadvertently affect the mycorrhizal communities found in association with the plants. A comprehensive study conducted on 3,000 plots with tens of thousands of trees across United States forest affected with nitrogen pollution was studied; plants showed symbiotic relationship with either ectomycorrhizal fungi which grow on surface or arbuscular mycorrhizal fungi which penetrate deep inside the roots. Lower abundance of ectomycorrhizal fungi in forests with high nitrogen pollution than trees having arbuscular mycorrhiza as their symbiotic partner was their major observation. This implies that nitrogen pollution caused by burning fossil fuels and excess usage of nitrogen based fertilizers on farms makes ectomycorrhizal fungi less abundant in forest (Averill *et al.*, 2018). Losing these fungal groups can increase decomposition and amount of carbon locked in soil and trees which can expedite the climate change process. Even if we keep climate change aside, several antibiotics are extracted from soil fungi and mushrooms that are ectomycorrhizal in nature and loss of ECM from forest means loss of such biologically important compounds from nature. Very few studies focus on these minor changes in nature that can ripple through the ecosystem and eventually they might change the functioning of that particular ecosystem. Interestingly a recent research by Bastin *et al.* (2019) proposes in their findings that planting trees in 350 million hectares could potentially reduce 26 Gt of atmospheric CO₂ that could prove beneficial in mitigating climate change.

Boreal peatlands cover approximately 17% of boreal zone and represent one of the largest storage of terrestrial carbon. Asemaninejad *et al.* (2018) assessed the effect of climate change on fungal communities and observed that the frequency of members of *Leotiomyces* was similar; however, *Sordariomyces* and *Eurotiomyces* increased after treatment. *Agaricomycetes*, *Chytridiomycota*, *Monoblepharidomycetes* and *Microbotryomycetes* increased with time and were a dominant group whereas an opposite trend was observed for *Zygomycota*. An interesting result of this study was that temperature, water content and organic content influenced fungal communities which in turn altered carbon dynamics of soil. Reasonable explanation given by the author is that during the early stage effect on water table was much pronounced as decomposers were more prominent while elevated temperatures altered the cell membrane composition. Genus *Mycena* and *Tricholoma* emerged as a dominant taxa at higher temperature; they are potential cellulose and lignocelluloses decomposer. Increase in *Ascomycota* group could be an important indication towards the establishment of vascular plants under high temperature in future.

Mucha *et al.* (2018) hypothesized that strong effects of global

warming prevail at biome boundary and northern latitudes. *Lactarius*, *Laccaria*, *Tomentella*, *Russula*, *Hebeloma* and *Tuber* are the most common ectomycorrhizal (ECM) taxa observed in boreal forest and *Laccaria*, *Russula*, *Clavulina*, *Tuber* and *Tomentella* responded at elevated temperature. *Lactarius* and *Russula* decreased in abundance whereas *Wilcoxina* and *Laccaria* increased at higher temperature while *Clavulina* was significantly more abundant in warmer temperature treatment. Therefore it might be predicted that the future climate change may favour more thermophilic and drought resistant micro-organisms.

A comparative study, for the first time, showed the long term effect of warming on the ECM fungal community in dry and moist tussock tundra region. *Cortinarius* is prevalent in the moist tundra region and *Tomentella* was found highly diverse in dry tundra region. Here was a significant decrease in the ECM fungi due to warming, however author suggests certain species are favoured by warming, while certain species go locally extinct due to direct and indirect effect of warming. There is a clear compositional shift in the fungal communities that might affect biogeochemical cycles and soil carbon sequestration (Morgado *et al.*, 2015). In an impressive study by Egidi *et al.* (2019), 235 soils collected globally from terrestrial ecosystems were analyzed for fungal dominance wherein *Ascomycota* was found to be dominant member all over the earth especially in tropical and temperate forests. The members of *Pezizomycotina*, *Leotiomyces*, *Eurotiomyces* and *Dothideomycetes*, *Tremellomycetes* and *Mucoromycotina* were ubiquitous in distribution. The dominance appeared to be linked to their higher stress tolerance and resource uptake, competitive capabilities like melanin deposition, as well as resistance to antibiotics and secretion of broad array of antibiotics. From the climate perspective this might be the group which is vulnerable as it exhibits habitat preference.

3.2. Marine ecosystem: A reasonable definition given by Pang *et al.* (2016) for marine fungus is “any fungus that is recovered repeatedly from marine habitats and: 1) is able to grow and/or sporulate (on substrata) in marine environments; 2) forms symbiotic relationships with other marine organisms; or 3) is shown to adapt and evolve at the genetic level or be metabolically active in marine environments.”

Nearly 70% part of planet earth is hydrosphere and dominated mainly by phototrophs and chemotrophs and together with other microbes constitute a population of around 10²⁹ (Flemming and Wuertz, 2019); fungal counts are around 10,000 species (Jones, 2011). Like terrestrial ecosystems, dominant fungal phyla include *Ascomycota* and *Basidiomycota* (Kohlmeyer and Kohlmeyer, 1979) including a high number of novel *Chytridiomycota* (Comeau *et al.*, 2016) which are closely related to terrestrial fungal species indicates a transitional switch to marine life (Schoch *et al.*, 2009). Marine ecosystems are important to study during climate change because marine phytoplanktons alone can fix ~50 Pg C per year and have faster turnover rates as compared to terrestrial plants (Behrenfeld, 2014), however turnover rates for marine fungi are yet unknown (Amend *et al.*, 2019). Mainly temperature, availability of nutrients and sunlight control the assembly of microbial community (Sunagawa *et*

al., 2015). Besides these three primary factors, water density, stratification, circulation, particulate organic matter, salinity fluctuations and organic or inorganic inputs by various water bodies in the ocean are also key deterministic factors (Amend *et al.*, 2019). Additionally marine environments are also influenced by changing climate that includes water acidification, shifts towards warmer temperature and anthropogenic pollution that indirectly puts marine life to threat. These factors regulate carbon and nitrogen biogeochemical cycles and thereby maintaining a balance between the nutrient input into the atmosphere, hence controlling the climate change (Hurd *et al.*, 2018).

That complex food webs are regulated by fungi in the marine environments has been well exemplified in the studies using DNA stable isotope probing DNA SIP with ^{13}C labeled diatom derived polysaccharide; *Cladosporium* directly assimilated phytoplankton derived organic matter (Cunliffe *et al.*, 2017). This clearly shows how one domain of life is dependent on the other for nutrition and survival. Therefore, slight perturbations in environmental conditions can shift food webs, primary productivity and carbon release from the oceans (Hutchins and Fu, 2017). Holding *et al.* (2015) have predicted that rising carbon dioxide levels in the atmosphere and in presence of surplus availability of nutrients in oceans can increase phytoplankton primary productivity. Also, global sea ice index has declined in past few years consequence of which light can penetrate deep into the oceans that will eventually increase the primary productivity (Kirchman *et al.*, 2009). However, Behrenfeld *et al.* (2017) predicted different patterns for productivity in polar regions and this somehow points the need for long term data in community composition. Marine ecosystems also face anthropogenic pollution in forms of plastic and limited studies point towards the capability of fungi to degrade them. *Aspergillus tubingensis*, a soil fungus (Khan *et al.*, 2017) and *Zalerion maritimum*, a marine fungus (Paco *et al.*, 2017) have the potential to degrade polyurethane and polyethylene to lessen down anthropogenic pollution such as plastic pollution, respectively.

Fungi also interact with other organisms and show mutualistic relationships that influence the overall health of the host. One such example is coral reefs, they are major sites vulnerable to climate change and fungi being the partner of the holobiont becomes an important area to study. Corals are strongly influenced by range of microbes in association (Torda *et al.*, 2017) and ocean acidification causes species specific difference in sensitivities towards changing environment (Comeau *et al.*, 2019). Perturbations in environmental cause decline in corals and possibly a shift in the ecosystem towards microalgae that is something certain to happen (Enochs *et al.*, 2015).

4. CARBON SEQUESTRATION

The amount of carbon dioxide released into the atmosphere as a result of decomposition of soil organic matter is called carbon footprint and this carbon dioxide emission has the potential to severe the climate change effects. Every act of decomposition process leaves a carbon footprint including microbes. The total carbon on the earth is balanced by the carbon cycle derived by mainly fungi and bacteria which

either act as generator or user of these gases in the cycle. As it is a proven fact that soil respiration increases with rise in temperature therefore the global warming is actually accelerating the metabolic processes of microbes and fluxing more green house gases rapidly into the atmosphere. Then the question arises how much do soil microbes contribute to atmospheric carbon dioxide? Indices such as carbon utilization efficiency (CUE) is the ratio of microbial biomass carbon to microbial biomass plus carbon dioxide, used to determine the efficiency of organic matter conversion to microbial products relative to carbon dioxide released via respiration (Sinsabaugh *et al.*, 2013). Global assessment of ecosystem carbon turnover time is found to be 23 years and carbon resides in soils near equator for a shorter time than at latitudes 75°North (Carvalhais *et al.*, 2014) indicating a clear dependence of turnover on temperature and precipitation. From the recent data the microbial contribution in release of carbon from the soil increased by 1.2% since 1900, surpassing plants' conversion of carbon dioxide to oxygen (Bailey *et al.*, 2018). Therefore, it is imperative to utilize those microbes that extract carbon dioxide from atmosphere and store it in soils (**Fig. 1**). Prof David Johnson of New Mexico State University vehemently recommended utilization of microbes with high fungal to bacterial ratio strongly towards fungi (unpublished data). His findings support the fact that increased ratio towards fungi effectively utilizes nutrients from soil and reduces the emission of carbon dioxide to the atmosphere (Berger, 2019).

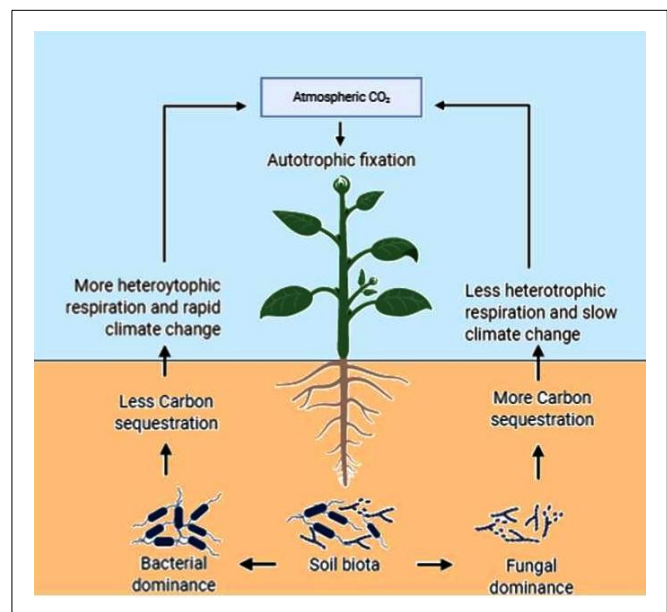


Fig. 1. Management of soil microbial communities to enhance carbon sequestration. Fungi are more efficient than bacteria in terms of abundance, high metabolic rate and are key determinant in deriving major biogeochemical cycles. They utilize root exudates released from plants fixed by photosynthesis for nutrition and lock them in soil as organic carbon and a fraction of which is released as CO_2 , CH_4 through the process of heterotrophic respiration and methanogenesis, respectively. Keeping a high fungal to bacterial ratio in soil can help to lock more carbon in soil, slow down CO_2 release and most likely to mitigate climate change.

5. FUNGI-SOIL-ATMOSPHERE CONTINUUM!

Climate has direct as well as indirect effects on fungal communities. Direct effect includes temperature and water that can affect fungi. Temperature rise can increase the abundance and activity of fungal species directly as they are favoured in warm climate. While the location lowered down the water table can compel fungi for adapting survival strategy using hyphae for long distance transport for water. On the other hand, indirect factors include plant communities that influence fungal partners residing as mutualists through photosynthetic exudates of roots (Treseder *et al.*, 2016).

5.1. Temperature: An interconnection exists between atmosphere and soil biodiversity compartments and for the first time an underground network of microbes that connects plant diversity has been mapped. A global survey extended in around 70 countries, gave a term 'wood wide web' to denote how the symbiotic association of fungi with plants alters with latitude. Warm and tropical forests near equator where decomposition is rapid are dominated by AMF whereas boreal forests near poles where decomposition is slow are predominantly occupied by ECM. What unique this study brings out is the remarkable difference in the diversity of AMF and ECM that changes with the change in the local climate forming transitional zones along the latitudes. This led authors to predict that as the earth warms, there is apparent shift of about 10% of EM associated trees to AM associated trees (Steidinger *et al.*, 2019). Staddon *et al.* (2002) have demonstrated an indirect effect on mycorrhizal communities due to temperature induced changes in plant communities. Any change in plant community due to climate change is reflected in the mycorrhizal communities. However, in contrast to this Newsham and colleagues (2016) using a pyrosequencing approach observed a direct relationship between the rising temperature and fungal diversity with an increase of 11.3 fungal taxa per degree Celsius rise in surface temperature. Warmer habitats favour higher diversity of fungi especially genus *Verrucaria* and among yeast *Rhodotorula* and *Helotiales*, *Cladosporium* and some unclassified OTU of ascomycete decrease in warmer regions. They deduced that there could be a rise of up to 27% in the diversity of fungi in Antarctic soils by 2100. There will be a significant increase in the nutrient decomposition hence more biological productivity. However, loss of trees in Amazonia can also result into loss of their symbiotic partners and as the long term 50 years record suggests these trees are fast declining due to elevated temperatures (Gloor, 2019). Replenishing the ecosystem with natural forests (Lewis *et al.*, 2019) and non native trees can improve the carbon sequestration from the atmosphere (Woodcock *et al.*, 2017).

5.2. Green house gases: The earth is surrounded by thick layer of gases and they trap heat from the atmosphere making the planet warmer than before. Such gases are termed as green house gases and those having predominant role in global warming are carbon dioxide, methane and nitrous oxide.

In a long term study of 26 years to deduce the soil and temperature relation it was observed that the world currently emits from fossil fuel burning about 10 billion metric tons of

carbon into the atmosphere in the form of carbon dioxide (Melillo *et al.*, 2017). The authors expect the situation to be more severe as at present soil organic matter in Arctic is locked in the ice but according to findings of Lee *et al.* (2017) arctic ice has started melting at a faster pace creating ice-free area and so the flux of carbon dioxide into the atmosphere produced upon decomposition of soil organic matter has risen. The severity of the matter could be well related from a recent study that reports that breakage of fast ice in Antarctica has led to disappearance of emperor penguin colony from Halley Bay site; scientists call it as a 'catastrophic breeding failure' since sea ice condition is a result of climate change (Fretwell and Trathan, 2019). Fungi are involved in storing carbon in soil, and soil with ECM can store 70% more carbon than soil without ECM. Therefore, one can argue that the amount of carbon stored and the rate at which it converts to atmospheric carbon dioxide is determined by ECM members. Staddon *et al.* (2002) demonstrated that elevated carbon dioxide only effects the plant growth and not the mycorrhizal partners of plants. A contrasting study by Tedersoo *et al.* (2014) states that climate change may potentially reduce ECM fungi thus expediting the process in the future. A study by Kohler *et al.* (2009) demonstrates the effect of carbon dioxide on fungal diversity in arid and semi arid Mediterranean regions. A negative trend of mycorrhizal colonization was observed with elevated carbon dioxide (760 ppm) and water stress and soil amendments with such beneficial strain could provide stability to other microbes under stressed conditions. In another study of similar nature negligible effect of carbon dioxide (757 ppm) was observed on fungal community structure and ITS gene copy number while there was increase in diversity of archaea and decrease in bacterial diversity. Bacterial communities are strongly determined by carbon dioxide concentration as compared to fungal communities, a striking dissimilarity was noted in this study. Fungal groups are slow growers and secondly labile carbon available was not enough to induce a shift that could result in the absence of response (Lee *et al.*, 2015).

Major source of nitrous oxide emission in agriculture soils is released though incomplete denitrification. A study utilizing AMF *Rhizophagus irregularis* to manage nitrous oxide production from *Zea mays* field demonstrates the importance of utilization of fungi in reducing the production of this potent green house gas (Storer *et al.*, 2018). Such studies suggest a future for green house gas reduction and climate change mitigation employing fungi.

5.3. Rainfall and drought: Fungal taxonomic shift was observed in a long term field monitoring on *Salix viminalis* (willow). There was around 30% decline in the *Basidiomycota* and significant decline in *Zygomycota* after a post extreme rainfall whereas least deviation in the *Ascomycota* group was observed. *Truncatella angustata*, *Plectopharella*, *Pilidium*, *Motierellaceae*, *Sporomiaceae*, *Cryptococcus* and lichenized fungi like *Venturiaceae* and *Verrucaria andesiatica* increased in abundance post rainfall (Barnes *et al.*, 2018). On the contrary in a study conducted for over a period of 17 years in grass land ecosystem, a remarkable difference was observed in summer drought condition on fungal assemblages wherein 66 of the 208 fungal

taxa were entirely absent. Shifts in fungi is related to leaf dry matter content, whereas changes in soil bacteria are related more to differences in plant C:N ratios (Sayer *et al.*, 2017). In a comparative analysis of fungal and bacterial communities in drought condition fungal evenness and richness increased during drought. Bacterial communities were not resilient as they were previously thought to be. Drought indicators belonged to the phyla *Ascomycota* and *Glomeromycota* and sensitive indicators included phylum *Zygomycota*. Moreover, unlike bacteria, fungal tolerant indicators were found to be more connected (de Vries *et al.*, 2018). Wet land response to climate change suggest that fungi are less sensitive than other microbial groups to external factors such as carbon dioxide and nitrogen (Lee *et al.*, 2015). Such studies are important in framing patterns of different ecosystems.

4.4. Wildfires: Climate is changing at a fast pace and with frequent wild fires of especially Colorado a new term has emerged, 'fire tsunami' (<https://grist.org/article/weve-entered-the-era-of-fire-tsunamis/>). Wildfires are a direct consequence of climate change and have become a serious threat to forest ecosystem as it impairs the recovery of endangered trees post wildfire. There are certain regions in western US which are assumed to surpass 'critical climate threshold,' a term that is used to indicate regions that may not return to normal after wildfires. Ponderosa pine and Douglas fir, important trees for timber industry are endangered and less likely to regenerate themselves after wildfire thus are on the verge of being locally extinct (Davis *et al.*, 2018). Fungi are an integral component of forest ecosystem and certain ECM are specific and grow in close association with only specific plant species, eg. *Russula delica* is more abundant on *Betula papyrifera* roots but is barely detected on *Quercus rubra* roots (Mucha *et al.*, 2018). A very limited studies suggest that these fungi could also become locally extinct in the face of climate change. Interestingly, Murata and colleagues (2017) in a study conducted on 104 Pine forests observed that *Rhizopogon* sp. associated with endangered *Pinus amamiana* forests was present at all sites that suggests a possible role of the species in seedling establishment of this tree and efforts could be made to conserve such endangered plants from facing extinction.

6. ADAPTATIONS IN FUNGI

There are now several reports that indicate that fungi adapt themselves to the changing environment. One such drug resistant fungus known as *Candida auris* has been reported and scientists have termed it as 'superfungus'. Scientists strongly believe that its ability to spread across the world is on account of its growth at warmer temperature (37°C) (Casadevall *et al.*, 2019). Sometimes adaptation leads to the advantage of reproductive success in mushroom assemblages that becomes significantly darker in areas with cold climate, an example of the adaptive selection residing in thermal environment. (Krah *et al.*, 2019). In a study by Andrew *et al.* (2016), authors observed the adaptation involving sporocarp production and spore volumes where smaller spore volumes were associated with warmer temperature and greater spore volumes were found to be associated with increased humidity. Species having veined hymenial layer like *Cantharellus*,

Craterellus and *Pseudocraterellus* were less abundant with increased humidity. It has also been noticed that increase in temperature can also affect the morphology of fungi through hyphal branching, radial extension rate and hyphal coverage (A'Bear *et al.*, 2012). Minor change in mean annual temperature by 0.2° C has an effect on timing of spore producing structure which can delay it by 1 day. Climate change can also have adverse effect on fungal reproductive timing (Andrew *et al.*, 2016). The increase in fungal diversity in warmer regions has its negative implication on human health as more heat tolerant fungi could emerge or they adapt to become thermally resistant (**Table 1**).

Table 1: Various adaptation in fungi due to climate change

| S. No. | Adaptation | Importance | Reference |
|--------|--------------------------|--|------------------------------|
| 1 | Thick walled spores | Thickening in walls of spores can help them withstand temperature and water stressed conditions | Koide <i>et al.</i> (2014) |
| 2 | Melanisation | Melanin serves dual purpose. Resist fungi from desiccation and may provide mechanical strength in case of drought to transport water from long distance through hyphae | Fernandez and Koide (2013) |
| 3 | Rhizomorphs | Specialized fungal structures to transport water and nutrients and to withstand harsh conditions of the environment | Yaffeto <i>et al.</i> (2009) |
| 4 | Jelly fungi | These fungi can withstand repeated heat and cold environment and can survive even under scarcity of water | Webster and Weber (2007) |
| 5 | Dormant spores/sclerotia | Sclerotia are dormant structures survive for several years in soil in dormant phase and can face harsh conditions like high temperatures and wildfire. Fungi may reappear under suitable conditions. | Koide <i>et al.</i> (2014) |

In a first long term study Romero Olivares *et al.* (2015) tested adaptation in strain of *Neurospora* at 16 °C and 28 °C. The mycelial growth rate and biomass was not altered whereas adapted strains produced more energetically expensive spores per unit biomass and higher mass specific respiration. Contrary to general expectations their findings did not support the idea that global warming will lead to increase in carbon use efficiency.

Another interesting example of how fungi respond to huge selection pressure exerted by climate change is observed in *Ophiocordyceps kimflemingiae* (zombie ant fungus) and its adaptation to local condition. This fungus infects carpenter ants without affecting their brains and while deriving nutrition from ants they also control the biting behaviour so that the tropical zombie ants always bite on leaves and temperate zombie ants always bites on twigs and bark (de Bekker *et al.*, 2015).

Lichens are highly vulnerable to climate change and there is sharp decline in their population in arctic / alpine regions in much of area now covered with vascular plants. They show diverse adaptive features in the climate change scenario. In drought condition they alter their morphology by increasing mass per area to retain more water during stressful condition. They also have the capability of switching photosynthetic partners or can also migrate (Larsson *et al.*, 2012).

7. CONCLUSION

Fungi are cosmopolitan in distribution and are efficient drivers of soil carbon feedback cycles than other microbial

species. They contribute towards a major portion of green house gases into the atmosphere. The situation is more alarming because “it implies we’ve added a lot of soil carbon back to the atmosphere as carbon dioxide, and our current models of the climate system do not represent this carbon dioxide emission pathway through fungi” (Averill *et al.*, 2018).

Laboratory experiments and data on mesocosm, *in situ* field experimentation and knowledge of fungal diversity and its evolutionary component required to understand any *in situ* change. Climate change demands long term study to observe effects that are important especially in nutrient poor quality soils and slow growing vegetation since many fungal species are slow growing therefore demand long term monitoring to understand the impacts of global climate change on fungal communities. Prominent shifts in community structure are reported in various studies. This knowledge could be utilized to introduce a new fungal genera or replenishing the existing fungal community like with AMF to successfully reduce green house gas emission or employing soil fungus *Trichoderma reesei* in biofuel production to lessen down the effect of climate change.

Key factors like temperature, rainfall, wildfires expedite the process of climate change due to which biodiversity is fast declining. However, due to lack of uncharacterized diversity of microbes, a connection between fungi and climate change is still unknown. As a result the native fungal species of any region might go locally extinct or might facilitate the establishment of invasive species that may pose competition or extinction of certain species across region sometimes which might therefore go undocumented. Above all, such studies suggest that anthropogenic signal for climate change is likely to increase in the near future and one might expect the fungal communities to eventually experience more substantial transformation in their community structure. Million of hectares of pristine tropical rain forest for paper and palm industry are destroyed in the past few years along with their ECM component and loss, if any, in such destruction remains undocumented. It alarms humankind to check such usages before consumption of plant derived products that confers an indirect effect on nature or climate. In culmination every action that mankind now makes requires its climatic ramifications to avoid major disturbances in nature.

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