University of Nevada, Reno

Developmental physiology, nitrogen preference, and estimated biofuel production of *Opuntia ficus-indica* (prickly pear cactus) in the United States

A dissertation submitted in partial fulfillment of the requirements for the degree of Doctor of Philosophy in Biochemistry

By

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Abstract

Opuntia ficus-indica (Prickly pear cactus) is a large succulent cactus species that has a long history of agricultural production for carmine dye, fruits, and edible young cladodes. As a CAM plant O. ficus-indica uses crassulacean acid metabolism (CAM) which is characterized by assimilation of CO_2 at night, which greatly increases water use efficiency. More recently, O. ficus-indica has been recognized as having high biomass and biogas (CH₄) production values comparable to other large CAM species. Opuntia eliator, a closely related tropical epiphyte species, has been shown to predominantly use C₃ photosynthesis in seedlings before switching to CAM photosynthesis. Similarly, young cladodes in *Opuntia ficus-indica* have been shown to predominantly fix CO₂ during the day until they mature. Some plants can also switch to CAM photosynthesis when preferred nitrogen sources (nitrate or ammonium) become limiting and O. ficus*indica* has been shown to have a slight increase in nitrogen uptake in ammonium vs.. nitrate. A field trial, and life cycle assessment has been conducted on biogas production from O. ficus-indica in Mexico, but biogas production values in the United States are unknown. Here, measured carbon isotope ratios, 24-hour gas exchange, and tissue acidity in O. ficus-indica seedlings and daughter cladodes to determine the occurrence of CAM vs. C₃ photosynthesis. Results demonstrated that O. ficus-indica seedlings used predominantly CAM photosynthesis to assimilated CO₂ even under well-watered conditions, while developing cladodes were shown to be sink tissues that switched directly from C₃ photosynthesis to CAM photosynthesis in greenhouse conditions. Here, N preference was

investigated by placing *O. ficus-indica* cladodes in sand culture and providing nutrient solutions with varying amounts of nitrate and ammonium for one month, and measuring differences in growth, biochemistry, and CAM and N-related gene expression. Statistical differences in *O. ficus-indica* growth, chlorophyll content, tissue acidity, soluble sugars, nitrate reductase activity, nitrate and ammonium content, glyoxylic acid content, N:C ratio and relative expression of genes involved N metabolism, and CAM activity were all detected when nitrate and ammonium were varied in sand culture. A life cycle inventory of biogas production from *O. ficus-indica* was built using results from an irrigation field trial that took place in Logandale, NV. Estimated biogas production from *O. ficus-indica* in the United States was 13,004.29-26,877.85 Nm³ ha⁻¹ yr⁻¹ produced from plants receiving 716 mm year⁻¹ which is comparable to that of actual production values in Mexico.

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Chapter 1

Introduction

State of our environment

Conveniences of the 21st century have made it easy to forget the connections between our environment and the commodities upon which we depend. The reality is that overall socio-economic performance is nested in the sustainable production of natural products from our ecological systems (Purvis et al., 2019). Ecological systems can be defined by the species composition at any point in time (Oliver et al., 2015), and human caused changes in climate, land use, and habitat quality can cause biodiversity loss to an extent that ecological systems might never return to prior equilibrium states.

Today, human activity is pushing the limits on nine proposed planetary boundaries (Steffen et al., 2015; O'Neill et al., 2018) that directly and indirectly have detrimental effects on ecological systems around the globe. These boundaries include climate change, biosphere integrity, land use changes, freshwater availability, biogeochemical flows (nitrogen and phosphorous cycles), ocean acidification, atmospheric aerosol loading, stratospheric ozone depletion, and the introduction of novel entities, such as invasive or genetically modified species, and chemicals with unknown effects. In the current state, high-risk boundaries of genetic diversity loss, phosphorus release, and nitrogen release have already been crossed.

Currently, global deserts, or regions where total surface water loss exceeds the surface water gain through precipitation, make up close to 40% of global land

area (Warner, 2009). Approximately 50% of the global land area is considered arid, semi-arid, or dry sub-humid (Zika and Erb, 2009). Desertification, or the process by which fertile grounds become desert due to deforestation, drought, and detrimental agricultural practices are now affecting an estimated one fifth of the global population (Geist, 2017). Global climate change in particular is projected to increase drought intensity by 30%, and result in a 5-fold increase in water demand in the 21st century (Naumann et al., 2018). In addition, climate change in combination with intense water use is causing a reduction in ground water stores (Cuthber et al., 2019). Agriculture accounts for 70% of global water demand (McDaniel et al., 2017), even though 95% of all crops are rainfed (Hadebe et al., 2017). Globally, 11% of croplands, and 10% of grass lands used for agriculture and livestock production are vulnerable to a reduction in water availability, particularly in Africa, Europe, and Asia (Fitton et al., 2019). In addition, each degree-Celsius increase in global mean temperature would also, on average, reduce global yields of wheat by 6.0%, rice by 3.2%, maize by 7.4%, and soybean by 3.1% (Zhao et al., 2017).

Plant photosynthesis

Of all the natural products that humans depend upon, those derived from agriculture are perhaps the most fundamental. Plants provide building materials for homes (Berge, 2007), fibers for textiles (Xu-fu, 2003), medicinal drug discoveries (Süntar, 2020), sources of bioenergy (Reid et al., 2020), food and fodder, and many other products that are economically valuable and crucial for a high quality of life. Additionally, plants play a role in maintaining global greenhouse gases, as land use changes associated with deforestation, logging, and modern cultivation practices caused a net CO₂ release of 1.5 ± 0.7 Gt of C yr⁻¹ into the atmosphere from 1990 to 2005 (Le Quéré et al., 2009). However, an estimated annual 10^{11} tons of global atmospheric carbon is also fixed by an enzyme known as D-ribulose-1,5-bisphosphate carboxylase/oxygenase (RUBISCO) by C₃ photosynthetic plant species (Hennacy and Jonikas, 2020). Thus, fixation by plant capture an estimated 25% of global carbon emissions

In traditional C₃ photosynthesis, RUBISCO combines ribulose-1,5bisphosphate with a single molecule of CO₂ to form a six-carbon molecule that is quickly hydrolyzed into two molecules of 3-phosphoglycerate (Taiz et al., 2015). The resulting 3-phophoglycerate molecules enter the Calvin-Benson-Bassham cycle, which ultimately regenerates more ribulose-1,5-bisphosphate to continue CO₂ assimilation by RUBISCO. Globally, an estimated 85% of all plants use C₃ photosynthesis, 5% use C₄ photosynthesis, and the remaining 10% use CAM photosynthesis (Kumar et al., 2017).

While most plants use C_3 photosynthesis, there are several trade-offs to using primarily RUBISCO for CO₂ assimilation. The first is that RUBISCO will occasionally fix O₂ instead of CO₂, which forms 2-phosphoglycolate (Busch, 2020). 2-Phosphoglycolate is converted into glyoxylate, which must be shuttled between the chloroplast, peroxisome, and mitochondria to be metabolized in an energetically costly process known as photorespiration. The second is that RUBISCO discriminates against the fixation of CO₂ containing C¹³ isotopes (O'Leary, 1988) except under water-limited conditions in which stomata are closed, in which case RUBISCO fixes all remaining CO₂ readily available (Leavitt et al., 2007). The third is the relative size of RUBISCO (560 kDa) and abundance of making up 30% of soluble protein and 25% of total leaf nitrogen (von Caemmerer and Quick, 2000), which requires a considerable N demand to maintain non-limiting RUBISCO concentrations. The fourth is the inherently slow catalytic turnover rate of about 3.5 s^{-1} per catalytic site at saturated CO₂ levels (Hennacy and Jonikas, 2020). Lastly, RUBISCO requires light for activation, and therefore, can only fix CO₂ during the day when sunlight is available.

Both C₄ photosynthetic and CAM pathways avoid the limitations associated with RUBISCO by first assimilating CO₂ into bicarbonate with carbonic anhydrase (CA) and then phospho*enol*pyruvate carboxylase (PEPC), which combines the resulting bicarbonate with phosphonyl-pyruvate (PEP) to form oxaloacetate (Cousins et al., 2007; Males and Griffiths, 2017). Compared with RUBISCO, CA/PEPC has a 60-fold higher reaction rate, readily fixes ¹²CO₂ and ¹³CO₂ isotopes, and does not fix O₂, suppressing limitations due to photorespiration (Lerman et al., 1974; Lüttge, 2002; Rodrigues et al., 2014). The C₄ photosynthetic pathway is characterized by a specialized interwoven anatomy and biochemistry that concentrates CO₂ into RUBISCO-containing cells eliminating photorespiration (Hatch, 1987). In C₄ photosynthesis, atmospheric CO₂ is fixed in mesophyll cells by CA/PEPC into malate. Malate is then shuttled into bundle sheath cells where it is decarboxylated. The decarboxylated CO₂ is then refixed by RUBISCO as in the normal C₃ photosynthetic pathway. Interestingly, the C₄ photosynthetic pathway is thought to have beneficial advantages under higher temperatures (Hennacy and Jonikas, 2020) where in C₃ photosynthetic plants CO₂ diffuses more readily out of the cell into large extracellular spaces, leaving a higher ratio of O₂ to be fixed by RUBISCO. In contrast, in C₄ plants, the carbon concentrating mechanisms in the bundle sheath cells limit photorespiration and improve water-use efficiency (WUE) defined as unit carbon gained per unit water lost.

Crassulacean acid metabolism

CAM photosynthesis also assimilates atmospheric CO₂ with PEPC, but unlike C₄ photosynthesis, the carboxylation and decarboxylation steps are temporally separated by a specialized diel circadian rhythm instead of a spatial separation made possible by specialized Krantz anatomy (Ting, 1985). Rather than opening stomata during the day as C₃ and C₄ photosynthetic plants do, obligate CAM plants keep stomata closed during all or part of the day, and open stomata at night. Nocturnal carbon assimilation is achieved by CA/PEPC, and fixed carbon is ultimately stored in the vacuole as malic acid until daybreak (Figure 1). During the day while stomata are closed, malic acid is transported back into the cytosol where it is decarboxylated by either cytosolic PEP carboxykinase (PEPCK) or NAD(P)-malic enzyme (ME) and freed CO₂ is refixed by RUBISCO in chloroplasts while sunlight is available (Silvera et al., 2010). The carboxylation, decarboxylation, and active transport of organic acids across vacuole membranes requires additional metabolic energy compared with C₃ photosynthesis (Lüttge, 2002). However, a contemporary computational model has demonstrated that this energy cost may be compensated by the carbonconcentrating mechanism of day time malate decarboxylation, especially under high light scenarios (Shameer et al., 2018). This diel circadian is often described taking place in four different phases, which are:

Phase 1: At (mid)night when stomata are open, CO₂ assimilation is occurring, and malate content is building up.

Phase 2: At dawn when the sun rises; and a small spike of CO_2 assimilation occurs by both CA/PEPC and light activated RUBISCO fixing CO_2

at the same time.

Phase 3: Sun is up, stomata are shut, and CO₂ is not being assimilated while malate content decreases with decarboxylation.

Phase 4: at dusk when the sun is setting, and a brief period of CO₂ assimilation occurs as the stomata open and both CA/PEPC and light activated

RUBISCO fixes CO₂ until light is no longer available.

Like C₄ photosynthetic plants, the main evolutionary advantage is an increase in WUE thanks to carbon-concentrating mechanisms and nocturnal CO₂ assimilation. By opening stomata at night, CAM plants are able to fix CO₂ with a 6-fold higher WUE than that of C₃ photosynthetic plants, and a 4-fold higher WUE than that of C₄ photosynthetic plants (Borland et al., 2009). C₄ photosynthesis and CAM are great examples of convergent evolution, as each are estimated to have evolved independently more than 60 times (Heyduk et al., 2019), and CAM in particular has been found in over 400 distinct genera across 36 families (Smith and Winter, 1996; Borland et al., 2009). Large, succulent cells are thought to be a prerequisite to CAM photosynthesis, as large tonoplast capacity is required for overnight storage of malate. The enlarged cells reduce mesophyll surface area that is exposed to intracellular airspace, reducing CO₂ conductance from the leaves compared to most C₃ plant species, although this trend is not observed in all CAM plant species.

Many coadapted traits are found in CAM plants that also enhance WUE and abiotic stress tolerance. Besides tissue succulence which can increase salinity tolerance and WUE, CAM plants can have dense epidermal trichomes, various water storage organs (bromeliad tanks, orchid pseudobulbs), thick cuticles and epidermal waxes, reduced stomatal density, enhanced stomatal responsiveness, and often rectifier like roots that pull away from the soil when dry to reduce water loss, and quickly return after rain (Niechayev et al., 2019; Pereira et al., 2021).

CAM species can also be separated by the type of decarboxylation enzymes and storage carbohydrates used (Christopher and Holtum, 1996, 1998). For decarboxylation, during the day malate that is released into the cytosol from the tonoplast is decarboxylated by either NAD(P)-malic enzyme (i.e., *Cactacea* and *Agave*), or NAD(P)-malate dehydrogenase (i.e., Bromeliaceae and Liliaceae). Plants that use NAD(P)-malic enzyme convert the resulting pyruvate into phospho*enol*pyruvate using pyruvate orthophosphate dikinase. Alternatively, some plants convert released malate into oxaloacetate with NAD(P)-malate dehydrogenase, and then oxaloacetate is decarboxylated with phospho*enol*pyruvate carboxykinase producing CO₂ and phospho*enol*pyruvate. Eudicot CAM plants typically store carbon as starch, whereas monocot CAM species store carbohydrates in the form of fructans; however, many variations on these general themes occur (Christopher and Holtum, 1996, 1998).

The occurrence of facultative CAM plants, or CAM plants that switch from C₃ photosynthesis to CAM under unfavorable conditions, has led to a further understanding in how C₃ plants might have evolved to perform CAM (Winter et al., 2015). The bimodal distribution of ¹³C abundance in several plant families (Silvera et al., 2010; Torres-Morales et al., 2020; Messerschmid et al., 2021) demonstrates that facultative CAM plants occur much less often than plants that use only C₃ photosynthesis or CAM with some exceptions. The occurrence of more C₃ photosynthesis or CAM species may be a result of niche availability strongly favoring one pathway or the other (Silvera et al., 2010; Torres-Morales et al., 2020). However, facultative CAM also provides advantages in regions that receive intermittent droughts as well (Leverett et al., 2021). Also, CAM plants have similar cellular anatomy to that of C₃ photosynthesis plants, and would not need to evolve new, specialized cell types to perform CAM as is the case with Krantz anatomy in C₄ photosynthesis species (Silvera et al., 2014). Indeed, evidence suggests that CAM photosynthetic species have been shown to have appeared in the fossil record long before C₄ photosynthetic species (Hennacy and Jonikas, 2020). The C₃ photosynthesis to CAM transition also requires CAM specific isozymes of enzymes that already occur in C₃ photosynthesis plants.

A developmental switch progression from C₃ photosynthesis to CAM has been observed in *Kalanchoe fedtschenkoi*, which initially performs C₃

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photosynthesis in distal young leaves, but the leaves switch to CAM once the leaves mature (Winter et al., 1982). In *Opuntia eliator*, a tropical epiphyte, seedlings predominantly perform C₃ photosynthesis, and assimilate the majority of diel CO₂ at night until subjected to water-deficit stress (Winter and Holtum, 2011). Likewise, very young cladodes of *O. ficus-indica* fix more net CO₂ during the light period than the dark period until subjected to water-deficit stress (Winter et al., 2008) even though *O. ficus-indica* is considered an obligate CAM specie. Developing cladodes have also been shown to be primarily sink tissues that receive carbohydrates from basal cladodes until they are large enough to begin CAM photosynthesis (Wang et al., 1998).

CAM plants that occur in especially arid environments, such as the cactus and agave families, have a higher occurrence of species that only use CAM photosynthesis and are thus commonly referred to as obligate CAM plants (Ehleringer and Osmond, 1989). Obligate CAM species often use CAM photosynthesis early on in development and continue to do so through maturity. Obligate CAM plants typically continue to assimilate CO₂ at night even when conditions become favorable. However, recent findings have shown that when conditions are optimum, the occurrence of CO₂ fixation at dusk and dawn by combined activity CA/PEPC and RUBISCO increases in obligate CAM plants to the point that net daytime assimilation may surpass net nighttime assimilation (Winter et al., 2008; Niechayev et al., 2019). Such findings indicate that while categorizing plants as C₃ photosynthesis, facultative and obligate CAM, the reality is that there is a continuum of CAM plasticity in many species (Silvera et al., 2010; Winter et al., 2015; Torres-Morales et al., 2020; Messerschmid et al., 2021; Winter and Smith, 2021).

Opuntia ficus-indica

One of the most emblematic, if not notorious obligate CAM plant species is *Opuntia stricta*. Introduced to Australia, it covered more than 25 million acres of land in 80 years and produced 1.5 billion tons of biomass (Osmond et al., 2008; Winter, 2021). Control with environmentally toxic herbicides failed and the cactus moth *Cactoblastis cactorum* proved to be an effective biological control. While highly invasive, *Opuntia ficus-indica* has been shown to have a negligible effect on plant species diversity by creating new favorable habitats for some species, but still altering plant species composition in the highlands of Eritrea (Tesfay and Kreyling, 2021). *Opuntia* spp. center of diversity is in central Mexico (Hernández-Pérez et al., 2009), and as a promiscuous and highly hybridizing plant species, there are currently 81 known varieties (Davis et al., 2019) with widely variable ploidy numbers (Palomino et al., 2016).

O. ficus-indica is a eudicot, platyopuntia (flat caldodes between nodes), terrestrial stem succulent with rectifying roots (North and Nobel, 1997), sunken stomata (Toumey, 1895), calcium druse crystals (Contreras-cladodeilla et al., 2011), spines, glochids (Mylo et al., 2021), and thick epidermal waxes and cuticles (Mayer, 2018). *O. ficus-indica* also produces tyrosine-derived betalin pigments instead of flavanol-derived anthocyanins (Smeriglio et al., 2019). Despite the common perception that CAM species are low yielding, *Opuntia* species have biomass productivity that vary widely from 2.4 to 47.3 Mg ha⁻¹ yr⁻¹ (Monjauze and Le Houérou, 1965; Garcia de Cortázar and Nobel, 1992; Dubeux et al., 2006; Cross et al., 2018; Ramírez-Arpide et al., 2018; Neupane et al., 2021). The upper end of this range exceeds yields of commodity crops that use C₃ or C₄ photosynthetic pathways; productivities of C₄ photosynthetic species grown for biofuels such as maize, switchgrass, and sugarcane range from 5 to 26 Mg ha⁻¹ yr⁻¹, and C₃ photosynthetic species grown for biofuels such as maize, species grown for biofuels such as oil palm, poplar, and willow produce between 2-14 Mg ha⁻¹ yr⁻¹ (Somerville et al., 2010).

From an economic perspective, *O. ficus-indica* has a rich and entertaining history. When the larva of the cochineal insect (*Dactylopius spp.*), generally considered a pest in modern day agricultural objectives (Mendel et al., 2020), which feeds on *O. ficus-indica*, and releases a deep red compound when crushed that is used to create the rich red dye known as carmine (Phipps, 2010). In the new world, cities conquered by Montezuma paid tribute in cochineal dye. Once the Spanish colonized the Americas, carmine found its way into thousands of paintings and was used to dye clothes all over the world. Spanish control of the carmine trade and production, most interestingly on the Canary Islands (Rowe, 2020), caused the value to become more expensive by weight than gold. Carmine was even used for the red coats in the British army, but only the officer's coats were made with carmine, while the regulars had to settle for dyes made with *Rubia tinctorum* (dyer's madder) (Phipps, 2010).

With the advent of synthetic dyes, the value of carmine diminished, but the global market for prickly pear fruits for beverages and jams, and young

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cladodes as vegetables remains today (Davis et al., 2019). More recently, the seed oil of *O. ficus-indica* fruits has been recognized as having a combination of unsaturated fatty acids including high amounts of linoleic acid (Ciriminna et al., 2017; Al-Naqeb et al., 2021), which is commonly found in expensive beauty products such as Argonne oil (Miklavčič et al., 2020). Each fruit contains anywhere between 50 to 300 large seeds.

Estimating Opuntia ficus-indica global productivity.

O. ficus-indica ecological physiology and agriculture have been studied for decades. Optimal growing conditions and tolerances are well known (Nobel, 2003). A systems dynamics model of CAM that accurately predicts diel CAM expression as an output (i.e., organic acid buildup, CO₂ assimilation) from measured biochemical and physiological inputs has been validated (Owen and Griffiths, 2013). Two separate mathematical models have been developed for estimating the potential productivity of O. ficus-indica in any given region. The first is an environmental productivity index (EPI) model first developed by (Nobel and Hartsock, 1984). This simple model estimates productivity using physiological response experiments that measure total nocturnal malate build-up due to changes in temperature, light, and water. Then, given average monthly climate data, the model can be used to calculate estimates of actual productivity in any given region. A newer, photo3 model, developed by (Hartzell et al., 2018) considers the soil-plant air continuum by incorporating the Farquhar model (Farquhar et al., 1980), and modifies it for a whole CAM plant by including CAM

circadian rhythm oscillator inputs, plant water storage, and atmospheric conditions; and considering soil-plant-atmosphere continuum as hydraulic fluxes in a convenient resistor-capacitor fashion.

The EPI model's strength is in its simplicity, as photosynthetic, hydraulic, and soil parameters required for the photo3 model are not necessary nor yet available for many CAM plants, although all inputs needed for O. ficus-indica are available and have been used in photo 3 predictions (Hartzell et al., 2018). The multiplicative index parameters make it easy to add parameters such as elevation (Nobel and Hartsock, 1986) and soil nutrients (Nobel, 1989) once those physiological responses are measured. However, the EPI model is based upon a monthly time scale and is limited in that it does not account for the interdependency between factors considered. For example, the PAR response of a given species might change when water is limiting or *vice versa* in ways not considered in the current EPI modeling approach. The Photo3 model, on the other hand, can make predictions on an hourly time scale, be used for C₃ and C₄ photosynthesis crop comparisons, and a comparison of EPI with Photo3 demonstrated that EPI and photo3 predictions can be up to 40% different (Hartzell et al., 2021), pointing out that EPI modeling is likely not the correct model to use when Photo3 estimation is possible.

The EPI model has been used in combination with GIS software to predict the range of *O. ficus-indica* with climate change (Owen and Griffiths, 2014), and the potential growing range is projected to expand in the southern United States. While there are projected range expansions, and *O. ficus-indica* biomass increases by 40% when CO₂ rates are doubled (Nobel and Israel, 1994), obligate CAM photosynthesis is mostly dependent upon nighttime temperatures (Neales, 1973). Slight increases in nighttime temperatures can reduce stomatal conductance of CAM species and reduce CO₂ assimilation, even under elevated atmospheric CO₂ conditions.

Nitrogen metabolism in Opuntia ficus-indica

While the EPI model developed by Nobel also includes a nutrient index (Nobel, 1989), the nutrient index is not species specific, and does not differentiate between nitrogen sources. Nitrogen metabolism is well understood in C_3 (Baslam et al., 2021) and C_4 photosynthesis plant species (Hirel et al., 2005). However, our current understanding of N metabolism in CAM plants is limited (Pereira and Cushman, 2019). Contemporary studies on CAM and nitrogen nutrition have focused on the link between CAM activity and nitrogen metabolism. Nitrogen assimilation into amino acids occurs in the chloroplasts via the glutamine synthetase and glutamine oxoglutarate aminotransferase (GS/GOGAT) cycle (Rodrigues et al., 2014). CAM plants placed in an increased atmospheric CO₂ concentration have an increase nitrogen uptake. Enzymes and substrates important for the tricarboxylic acid cycle and CAM photosynthesis are also important for nitrogen metabolism and amino acid synthesis. For instance, PEPC has been demonstrated to be important for ammonium assimilation and is directly regulated by quantities of 2-oxoglutarate and citrate synthase activity, both are important for amino acid synthesis. Malic enzyme (ME), and pyruvate orthophosphate dikinase

(PPDK), enzymes responsible for malic acid decarboxylation and phopho*enol*pyruvate (PEP) regeneration, respectively, are also important for providing carbon skeletons and metabolic energy for amino acid synthesis.

Contemporary studies on CAM and nitrogen nutrition have focused on the link between CAM activity and nitrogen metabolism. Nitrogen assimilation into amino acids occurs in the chloroplasts via the GS/GOGAT cycle (Rodrigues et al., 2014; Pereira and Cushman, 2019).

As is true for all plant species, variations in the preference of nitrogen source are known, whether it is ammonium, nitrate, or urea (Habteselassie et al., 2013; Wan et al., 2014; Hachiya and Sakakibara, 2017). In K. blossfeldiana, CAM expression has been shown to increase when plants are given more nitrate than ammonium, and when both nitrate and ammonium are absent or limiting (1) mM) (Ota, 1988). In Kalanchoe spp., CAM expression decreases when both ammonium and nitrate are supplied in high abundance (>5 mM), both together and separately presumably due to ammonium and nitrate toxicities (Pereira et al., 2017). Proton transport in tonoplasts mediating nocturnal organic acid storage in K. laxiflora and K. delagoensis is influenced by inorganic nitrogen sources, wherein low nitrate concentration increased nocturnal organic-acid accumulation, whereas the opposite was true in high nitrate concentrations (Pereira et al., 2017). The authors explain that this is likely due to an inhibition of tonoplast ATPase with high nitrate concentrations that corresponds to an increase in anionic species, which requires extra active proton pumping to maintain a beneficial electrochemical gradient. Collectively, these results demonstrate that in

Kalanchoe spp., CAM is highly expressed when plants are cultivated on soils with nitrate rather than ammonium, and when nitrogen in general is limiting, although the exact favorable amounts and preferences for ammonium *versus* nitrate varies from species to species (Hachiya and Sakakibara, 2017; Islam et al., 2020)

Several studies have documented the productivity of *O. ficus-indica* under different fertilizer treatments in the field (Garcia de Cortázar and Nobel, 1992; Dubeux et al., 2006; Sánchez et al., 2012; Nkoi et al., 2021) and the commercial N input is between 50–100 kg ha–1 year–1 (Davis et al., 2019). However, these studies were conducted with a wide variety of N sources, soil types, and production goals (*e.g.*, fruit, cladodes, seeds, and methane production). In one study, *O. ficus-indica* initially took up more N when given ammonium than nitrate, and therefore, *O. ficus-indica* likely has a slight preference for ammonium, but can fix either form of N (Vázquez et al., 2000).

Opuntia ficus-indica as a biofuel feedstock

With the addition of aerobic pre-treatments, methane yields from bioenergy crops can be increased. Nine-hours of aerobic pre-treatment of *O. ficusindica* cladodes using cow rumen fluids caused methane yields to increase 123% with a maximum yield of 0.72 L/kg volatile solid (Myovela et al., 2019). Further studies need to be conducted to assess the economic viability of *Opuntia* spp. as a biogas resource, but initial estimates seem to hold great promise. For example, annual electrical power generated globally from methane gas (about 5 PW) could be generated from CAM plants (e.g., *O. ficus-indica*) on 100–380 Mha of semi-

arid land or between 4% and 15% of such currently available land area (Mason et al. 2015). An evaluation of methane production from representative species of five different CAM genera (e.g., Ananas comosus, Agave angustifolia, Euphorbia virosa, Kalanchoe daigremontiana, and Opuntia fragilis) showed that A. angustifolia produced the highest methane yields and that biomethane yields were similar to those obtained from maize biomass (Lueangwattanapong et al., 2020). The CAM species *Euphorbia tirucalli* and *O. ficus-indica* are able to produce 1791 m³ and 1860 m³ of methane from 1 ha of land planted at high planting densities, respectively (Krümpel et al., 2020). Addition of microbes found in the soil of an O. ficus-indica plantation, in particular, Pectobacterium cacticida, and Enterobacter sp., were shown to digest O. ficus indica cladodes and have pectinase activity (Blair et al., 2021). These studies suggest that low-lignin CAM species can serve as highly attractive bioenergy feedstocks for biorefinery industries. The highest reported cellulosic ethanol production value from O. ficus*indica* is 2.6% v/w (Kuloyo et al., 2014), which is much lower than Agave *tequiliana*, as the higher total soluble solid fraction in juice content, and higher total dry mass make A. tequilana more suitable for ethanol production (Yang et al., 2015).

Estimating global impact of *Opuntia ficus-indica* production pathways

The environmental impact of a crop can vary depending upon the pollutant release associated with required land-use changes, conversion efficiencies to desired products, transportation, and other factors. Life cycle assessment (LCA)

models are a common tool for assessing the overall emissions produced from initial planting to final production and transport to market (Yan et al., 2011). In order to be able to compare the results of LCA results from different crops, most LCA calculations follow a standardized frame work (ISO) in what goal and scope definitions are first defined, second a life cycle inventory of all agricultural production inputs and outputs, then a life cycle impact assessment that determines the net emission produced per unit energy generated, and finally an interpretation of the results. A complete LCA of Opuntia-ficus indica biogas production with cow manure has been conducted using the results of actual field production values in Mexico (Ramírez-Arpide et al., 2018). The authors tested a scenario in which convention farming techniques were used with synthetic fertilizer, and an organic farming method, in which cow manure was used as fertilizer had energy return on investment values of 8.1 to 12.4 respectively. They found that the organic farming system decreased the overall environmental impact, but increased the released acids, and eutrophication (nitrogen pollution) on the field site. Overall, there was a 2.3% reduction in global warming potential and a 1.7% reduction in photochemical ozone creation potential (Ramírez-Arpide et al., 2018; Ramírez-Arpide et al., 2018).

With increased drought and temperatures associated with climate change, *O. ficus-indica* appears to have substantial potential as a fuel, feed, fodder and secondary products in the United States. Currently, collective *O. ficus-indica* products are worth an estimated \$2520 million year⁻¹ globally (Davis et al., 2019). *O. ficus-indica* growing operations could be combined photovoltaic arrays where water used to clean the panels also irrigates plants (Cushman et al., 2015), and reduce greenhouse gas emissions and generation of ozone deteriorating pollutants while creating a new source of energy (Ramírez-Arpide et al., 2018).

Developmental physiology, nitrogen preference, and estimates of biofuel production in *Opuntia ficus-indica*

This dissertation is divided into five independent chapters. The second investigates whether or not C₃ photosynthesis is used in developing seedlings and daughter cladodes before switching to CAM photosynthesis, the third investigates responses of *O. ficus-indica* growth, biochemistry, and expression of CAM and nitrogen-related genes under variation in provided nitrate and ammonium concentrations in sand culture, and the fourth chapter outlines ongoing life cycle assessment efforts of theoretical biogas and bioethanol production from *O. ficus-indica* using ground-truth results from a long-term irrigation trial conducted in the United States. These three studies provide novel insights into the relative contribution of C₃ photosynthesis to the high productivity observed for *O. ficus-indica* productivity while limiting run-off of unused N, estimations of biogas and bioethanol production in the United States, and lastly, a final chapter that provides a brief synthesis of the findings.

Figures



Figure 1: A simplified model of CAM with the black arrows representing fluxes at night, and the yellow arrows representing fluxes during the day. CO₂ entering the cell through nocturnally open stomata is converted into bicarbonate by carbonic anhydrase (CA) followed by fixation into phospho*enol*pyruvate (PEP) into oxaloacetate (OAA) by a CAM-specific phosphoenolpyruvate carboxylase (PEPC) isozyme (Nimmo, 2000; Boxall et al., 2017). This PEPC is constitutively expressed with its malate inhibition relieved through phosphorylation by a dedicated, circadian controlled phosphoenolpyruvate carboxylase kinase (PPCK) (Hartwell et al., 1999; Taybi et al., 2000). The resulting OAA is converted malate by malate dehydrogenase and shuttled into the vacuole by a putative aluminum malate transporter (ALMT) (Borland et al., 2016; Pereira et al., 2017) for storage until sunlight is available. During the day, stored malate is released from the vacuole through a tonoplast decarboxylation transporter (TDT), and

decarboxylated by NAD(P)-malic enzyme (Winter and Smith, 2021). The released CO₂ is then fixed by the Calvin-Benson-Bassham cycle by ribulose-1,5bisphosphate carboxylase/oxygenase (RUBISCO) as in the C₃ photosynthetic pathway. All chemical structures were sourced from chemspider.com.

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Chapter 2

Developmental Dynamics of Crassulacean Acid Metabolism (CAM) in *Opuntia ficus-indica*

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Abstract

Large agricultural species of cactus pear (Opuntia ficus-indica) show great potential for food, feed, and biofuel production in semi-arid and arid regions across the globe. However, the developmental basis of O. ficus-indica productivity in terms of its use of C₃ photosynthesis and crassulacean acid metabolism (CAM) remains poorly understood. Utilization of the C₃ photosynthetic pathway early in development might explain the high productivity of this obligate CAM species. The developmental progression of CAM was assessed in seedlings and cladodes developing on mature plants by titratable acidity, δ^{13} C carbon isotopic ratios, and daily gas exchange measurements. Titratable acidity of seedlings revealed a significant buildup of nocturnal tissue acidity in cladodes and cotyledons 75 days after germination. Fifty-day-old seedlings showed δ^{13} C ‰ values of -18.15 +/-0.31 typical of weak CAM plants. CO₂ assimilation occurred at dusk and night (phases 1 and 4) in seedlings, with predominant net CO₂ assimilation occurring at night once the primary cladode reached 5 cm in size. Slightly higher titratable acidity and δ^{13} C ‰ values

indicative of CAM was observed in seedling cladodes compared to cotyledons. In daughter cladodes, nocturnal titratable acidity build-up began to increase when cladodes were approximately 5 cm in height and peaked at 20 cm in height. Isotopic mass spectrometric analysis revealed δ^{13} C ‰ mean values between -14.77 ‰ and -15.30 ‰ (typical of CAM plants) regardless of cladode size. Daily 24-hour gas exchange measurements showed that the net daily CO₂ uptake was negative until cladodes were greater than 10 cm in length and that CO₂ uptake occurs primarily at night under greenhouse conditions. Collectively, these results suggest that developing *O. ficus-indica* cladodes begin as respiring carbon sink tissues that do not perform CAM and then begin to use CAM once cladodes reach 5 cm in length. Overall, these results demonstrate that CAM photosynthesis is the dominant form of carbon assimilation for *O. ficus-indica* even at the earliest stages of development.

Introduction

Crassulacean acid metabolism is an alternative mode of photosynthesis in which stomata open at night for carbon uptake and transpiration instead of during the day, as is the case in C_3 and C_4 photosynthetic plants. CAM results in a fourto six-fold increase water-use efficiency (WUE) relative to C_4 and C_3 photosynthetic species, respectively (Borland et al., 2009). CAM plants use carbonic anhydrase (CA) and phospho*enol*pyruvate carboxylase (PEPC) for the initial fixation of CO_2 at night in addition to ribulose-1,5-bisphosphate carboxylase-oxygenase (RUBISCO), which has a much lower reaction rate, is a larger enzyme, discriminates against CO₂ containing C¹³ isotopes, and occasionally fixes O₂ forming 2-phosphoglycolate, a product that must be regenerated back to 3-phosphoglycerate using photorespiration, an energetically wasteful process, before reentering the Calvin-Benson-Bassham cycle. Along with superior WUE, the carbon concentrating mechanism of CAM confers higher sun fleck capture efficiency to *Aechmea magdalenae*, which occupies lower canopy, light-limited environments (Skillman and Winter, 1997). CAM also confers a competitive advantage to plants occupying habitats where daytime ambient CO₂ levels and soil water may be limited as has been shown for the semiaquatic *Isoetes* genus (Keeley, 1981, 1983; Pedersen et al., 2011; Yang and Liu, 2015).

Some plant species also occur as C₃ photosynthesis-CAM intermediates known as facultative CAM plant species (Cushman and Bohnert, 1999; Borland et al., 2011; Winter et al., 2015). Several facultative CAM species perform C₃ photosynthesis, CAM, or switch between C₃ photosynthesis and CAM in response to salinity (Bohnert et al., 2001), temperature (Nievola et al., 2005), nutrient deficiencies (Rodrigues et al., 2014) including changes in N source (Pereira et al., 2017), and changes in osmotic potentials (Cushman and Borland, 2002). Facultative CAM species are thought to exist as an evolutionary transitionary state between C₃ photosynthetic and CAM as they tend to occur at lower frequencies than plant species using only C₃ photosynthesis or CAM (Silvera et al., 2010; Winter et al., 2015; Heyduk et al., 2019). However, the ability to switch between C₃ photosynthesis and CAM photosynthesis also provides fitness advantages in environments where daily water input is unpredictable (Cushman and Bohnert, 1999), or in regions that experience only brief periods of seasonal drought (Leverett et al., 2021). The transition from C₃ photosynthesis to CAM can also be developmental (Winter et al., 2011). For example, the first leaflets of a growing *Kalanchoe fedtschenkoi* perform C₃ photosynthesis, whereas more mature leaves transition developmentally to CAM (Winter et al., 1982).

In past studies, dawn/dusk tissue acidity, C¹²:C¹³ isotope ratio, and diel gas exchange measurements have been used to survey the amount of CAM occurring across a multitude of plant species (Messerschmid et al., 2021). Nocturnal stomatal conductance, fixation of CO₂ by PEPC, and accumulation and decarboxylation cycles of malate are all traits not unique to CAM plant species (Winter et al., 2015). However, the linked nocturnal acid accumulation with PEPC carbon assimilation is indicative of CAM (Ting, 1985) and has been used as a proxy of productivity for CAM plant species by measuring the changes in dawn to dusk acidity build-up with changes in environment (Nobel and Hartsock, 1983; Nobel, 2010). Fixation of CO₂ containing ¹²C and ¹³C isotopes by PEPC in CAM vs. mostly CO_2 containing ¹²C fixed by RUBISCO in the C₃ photosynthetic pathway means that the measured ratio of ¹²C:¹³C in a tissue sample can be used to reveal how much CAM activity was occurring over the lifetime of the plant (O'Leary, 1988; Silvera et al., 2010). Lastly, diel gas exchange measurements can be performed to the rate and duration of CO₂ assimilation over the course of a whole day (Chomthong and Griffiths, 2020).

Diel CO₂ uptake in CAM plants is often described in four separate phases over a period of 24 hours (Dittrich et al., 1973; Osmond, 1978; Owen and Griffiths, 2013; Chomthong and Griffiths, 2020). Phase 1 begins in the dark when stomata are open, and CO₂ is fixed by CA and PEPC into a 4-carbon acid, usually malate, that is ultimately stored in the vacuole overnight. Phase 2 happens as the sun begins to rise, and stomata are still open, resulting in a sharp spike of CO₂ fixation by combined activities of both CA/PEPC and light-activated RuBisCO. Phase 3 occurs mid-day when there is full sunlight, and net CO₂ assimilation is generally naught as stomata are shut during this time. During phase 3, malate is released into the mesophyll cytosol, and is decarboxylated by NAD(P)-malic enzyme into pyruvate and CO₂, which both enter the Calvin-Benson-Bassham cycle as in the normal C₃ photosynthetic pathway. Lastly, phase 4 begins at dusk, with another spike of CO₂ fixation as the stomata begin to open due to a decline in mesophyll CO₂, and CA/PEPC and RUBISCO both fix carbon as the sun sets.

CAM plant species that are adapted to xeric regions, such as *Opuntia ficus-indica*, are often described as obligate or constitutive CAM plants, which only utilize CAM in mature tissues for the duration of their life cycle, and under all abiotic conditions (Kluge and Ting, 1978; Osmond, 1978; Nobel, 1988; Winter et al., 2008). Cactus pear (*O. ficus-indica*) is a large photosynthetic stem succulent with large flat cladodes between each node (a platyopuntia *spp*.) with a rich agricultural history and center of diversity in central and southern Mexico (Griffith, 2004; Majur et al., 2012). The combined high growth rate, WUE, clonal propagation, and interspecific promiscuous breeding with other *Opuntia* cultivars has contributed to its success for the agricultural production of edible cladodes, fruits, and fodder, and as a noxious weed in arid and semi-arid regions around the globe (DeFelice, 2004). More recently, *O. ficus-indica* has been recognized for methane (Mason et al., 2015; Krümpel et al., 2020) and bioethanol (Kuloyo et al., 2014) production, and cosmetic and cooking oil production from over 200 large seeds per fruit (El Mannoubi et al., 2009; Ciriminna et al., 2017).

Investigation of the use of C₃ photosynthesis *vs*. CAM in *Opuntia elatior*, a tropical epiphyte related to *O. ficus-indica* revealed that seedlings fixed the majority of CO₂ during the day, and only fixed more CO₂ at night when the plants were exposed to drought (Winter and Holtum, 2011). When the cotyledons initially emerged, 24-hour gas exchange was similar to that of C₃ photosynthesis plants in that all carbon was fixed during the day, and none of the phases of CAM were evident. As the central cladode emerged, seedling gas exchange patterns began to show all four phases of CAM, but CO₂ was predominantly fixed during phase 2 and 3, when light was available. Titratable acidity also increased with age, but more so with drought. *O. ficus-indica* seeds have been germinated under laboratory conditions (Altare et al., 2006; Khan, 2006; Podda et al., 2017; Mokotjomela et al., 2021). However, the degree of C₃ photosynthesis *vs*. CAM in the desert adapted *O. ficus-indica* seedlings has yet to be investigated.

Changes in *O. ficus-indica* cladode diel CO₂ uptake have been tested with changes in light (Nobel and Hartsock, 1983), temperature (Nobel and Hartsock, 1984), addition of gibberellic acid (De la Barrera and Nobel, 2004), fruit production (Nobel, 2000), in mother (base) cladodes with and without daughter

(distal) cladodes under water-deficit stress conditions, under shaded conditions (Pimienta-Barrios et al., 2007), under elevated CO_2 and light levels (Cui and Nobel, 1994; Nobel and Israel, 1994; North et al., 1995), soil volume (Nobel et al., 1994), detached cladodes (Raveh and Nobel, 1999), and in one- and twoyear-old cladodes on fully grown plants in the field (Liguori et al., 2013). In general, O. ficus-indica diel gas exchange primarily occurs in phase 1 with the only change being a reduction in the rate and duration of gas exchange and nocturnal acidity build-up when conditions are unfavorable. Brief periods of CO₂ assimilation during phase 3 and 4 have been recorded in favorable conditions, but sharp spikes of CO₂ assimilation in phase 2 and 4 have generally not been observed in this species. O. ficus-indica cladodes begin as sink tissues when they emerge and become source tissues about 27 days after emergence (Wang et al., 1998). As the daughter cladodes grow, they initially show more CO_2 respiration than assimilation, an increase in carbohydrate-metabolism enzyme activities related to sink tissues at day 14, a gradual increase in nocturnal malate build-up, and initially import more carbon from mother cladodes. Interestingly, PEPC and RUBISCO activities were found to decrease as the cladodes grew, but this relationship was explained as being tied with decreasing cellular pH. Young cladodes (referred to as very young) from O. ficus-indica showed an increase in net CO_2 uptake in both the light and dark periods at a very similar magnitude as the cladode grew, and only when cladodes were water-deficit stressed did they show that net nocturnal CO₂ uptake overtook net daily CO₂ uptake (Winter et al., 2008). Net CO₂ uptake in the light increased when cladodes were re-watered, but

never was equal to or greater than nocturnal CO_2 uptake again after re-watering. The authors concluded that categorization of facultative and constitutive CAM may be convenient, but the reality is that CAM expression in plants is regulated on a continuum and not necessarily along strict categories of activity.

In this study, we investigated the occurrence of C_3 photosynthesis and CAM as *O. ficus-indica* seedlings and developing daughter cladodes grow by measuring dawn/dusk tissue acidity, ¹²C and ¹³C isotopes, and 24-hour gas exchange. The objective was to determine if the obligate CAM species *O. ficus-indica* only uses CAM photosynthesis in developing tissues, or if C_3 photosynthesis plays a part in the high growth rate observed in this species.

Methods

Opuntia ficus-indica seed collection, cleaning, and germination

To produce seeds for experimentation, four-year-old *O. ficus-indica* plants were transferred from 11.3 l pots to 50 l pots in 4:1 ratio of Sunshine MVP soil mix (Sun Gro Horticulture, Agawam, MA) and natural play sand (SAKRETE, Charlotte NC) in the University of Nevada, Reno valley road greenhouse complex. Plants were given 200 g of Osmocote (N-P-K of 14-14-14) as fertilizer and 8 grams of systemic insecticide treatment (Marathon[®] 1% Granular, OHP) every 6 months to control for thrips. Standard greenhouse conditions with natural light were approx. 1,100-1,500 µmol m⁻² s⁻¹ and temperature ranged from 28-32 °C day/17-18 °C night. In May, flowers were cross pollinated using cotton swabs. Over the proceeding months, fully mature fruits were collected for seed extraction. Seed extraction was performed by first removing the peel with a razor blade. The excess pulp was separated by hand, and the remaining seed/pulp mix was placed in a steel mesh colander (2133, Norpro 3 1/4 inch) and rinsed. Seeds were then placed in 500 ml beakers (one per fruit) filled with distilled water at room temperature with a stir bar. Seeds were rinsed in the colander and placed back in beakers with fresh distilled water daily over a period of several days until all of the pectin had been removed from the seeds. At this point, seeds were folded in a paper towel, placed in a drawer to dry, and then placed into appropriately labeled 50 ml Falcon tubes (352070, Corning). Cleaned seeds were allowed to age at least two years before germination.

Prior to planting, seeds were sterilized in a solution containing 10% v/v bleach and 0.5% v/v Triton X-100. Seeds were placed in this sterilizing solution with at least 10 ml of solution per seed. The seeds remained in the sterilizing solution for at least 10 minutes with stirring. After sterilization, seeds were moved into a laminar flow hood while still in solution and poured into a sterilized steel colander. Seeds were then washed with three rounds of sterile, autoclaved water, one round of 200 proof ethanol (V1016, Koptec), and allowed to dry.

After drying, seeds were clipped with sterile nail clippers (Revlon) to allow the germination media to make contact with the embryo. Nail clippers were used to carefully remove the seedcoat from the radicle side of the seed and along the outer edge while avoiding crushing or cutting of the embryo (Figure 1B). Two to three clips were necessary to give enough room for the emerging seedling to emerge from the seed coat. Clipped seeds were then submerged in germination media containing either 3.0% w/v sucrose, 0.6% w/v Phyto agar (PTP01-1KG, Caisson Labs), 0.5% w/v Murashige & Skoog medium with Gamborg's B5 vitamins (M404, PytoTechnology Laboratories), and 10 mM gibberellic acid (G-120-50, Gold Biotechnology Inc.) or the same media containing no sucrose in 25 x 100 mm sterile polycarbonate Petri dishes (89107-632, VWR). Petri dishes containing clipped seeds in germination media were then placed in a tissue culture growth chamber (AR-7522, Perceival Scientific, Inc. Derry. IA) set at 16/8 day/night cycles held at 24 °C and ~80 μ mol m⁻² s⁻¹ provided by fluorescent lamps. The percentage and timing of germination was recorded. After a two-week period, germinated seedlings were either transferred into plates containing 3.0% w/v sucrose, 0.6% w/v Phytoagar, 0.5% Murashige & Skoog medium with Gamborg's B5 vitamins, and no gibberellic acid, or the same media without sucrose.

Titratable acidity and isotopic mass spectrometric analysis of *O. ficus-indica* seedlings.

O. ficus-indica seedlings were destructively sampled after 30-days, 50days, 75-days, and 100-days post germination for tissue acidity and carbon isotope measurements. For both measurements, seedlings were harvested and separated into cladode and cotyledon tissues, which were then weighed. For tissue acidity, seedlings were harvested at dusk and dawn and immediately flash frozen in liquid N₂. The acidity was then analyzed using a modified protocol from Gehrig et al., 2005. Briefly, samples were placed in 25 ml beakers with 20 ml of 50% v/v methanol and heated until 10 ml volume was lost to ensure that all methanol was boiled off, and that the water reached a minimum of 80 °C. Boiled samples were cooled to room temperature and brought back to 10 ml with deionized water. For each sample the initial pH was recorded and then titrated to a pH = 7 for malate equivalent, 8.4 and 10 in case the previous pH points were missed. All samples were titrated with 5 mM KOH.

The H⁺ equivalent was calculated using the equation:

$$H = \nu * \left(\frac{0.01}{w}\right) * 1000$$

Where: $H = H^+$ equivalent, v = volume at pH in consideration, and w = fresh weight of the sample.

The difference between the average H^+ equivalent at dusk and at dawn was equal to the amount of malate (pH = 7) built up over night.

Samples collected for isotopic spectrometric analysis were collected at noon on the correct age after germination described above, dried for 72 hours in a LabConCo Freeze Dry System (Freezone 18), and pulverized using a Benchmark D2400 BeadBlaster Microtube Homogenizer. Between two and three mg of tissue was then loaded into tins (041070, Costech Analytical Technologies Inc., Valencia, CA), placed into a 96-well plate, and shipped to either the Facility for Isotope Ratio Mass Spectrometry (FIRMS) located at the University of California, Riverside (https://ccb.ucr.edu/facilities/firms) or the UC Davis, Stable Isotope Facility (SIF) at University of California, Davis (http://icpms.ucdavis.edu/facilities-procedures) for the analysis of the ratio of ¹²C and ¹³C isotopes in each sample using Vienna Pee Dee Belemnite (VPDB) as the reference standard.

Seedling gas exchange measurements

After germination, and an approximate one-month period in media that did not contain GA, seedlings with fully emerged cotyledons were transferred to 65 mm LI-COR pots (610-09646, LI-COR Environmental) with autoclaved Sunshine MVP soil mix (Sun Gro Horticulture, Agawam, MA) for gas-exchange experiments. Once the seedlings were potted, the entire pot with plant was placed in a plastic container with about 3 cm deep sterilized water to allow the soil to take up water through the drainage holes and maintain field capacity without changing the soil level or physical structure (Figure 1C). Plants were kept on a custom-built fluorescent light rack and moved closer or farther away from the light source to maintain either a PAR of 100 or 300 µmol m⁻² s⁻¹ depending on the experiment. The dates that seeds were placed in germination media, transferred to regular media, transferred to soil, and measured in the LI-COR portable photosynthesis system were all recorded. Just prior to gas exchange measurements, size of cotyledons, central cladode, and stem height were measured, and several pictures were taken from both side and top views.

In the first experiment, gas exchange was recorded over a 48-hour period in two separate 24-hour measurements using the LI-COR 6400XT portable photosynthesis system with the whole plant *Arabidopsis* chamber with clear top attached. In the first measurement, the LI-COR mixer was set to hold the sample chamber CO₂ at 400 μ mol mol⁻¹, the flow rate was set to 500 μ mol s⁻¹, leaf fan to fast (setting 5), block temperature was set to 20 °C, and the photosynthetically active radiation (PAR) at plant level was 100 μ mol m⁻² s⁻¹ with a 12-hour day/night photoperiod. The auto program (auto log 2) was set to log photo once every 5 minutes for 1440 minutes and match after every log. Desiccant (2088701, LI-COR) and soda lime (9964-090, LI-COR Biosciences) were changed every 3-4 hours. Humidity was held between 50-70% within the sample period. Interreference from root respiration was avoided by using a LI-COR 6400 exhaust set up that forces airflow down through the soil by restricting 50% percent of the exhaust flow (Figure 2 in:

https://www.licor.com/documents/89lw0pf0ooypx3wgfmnr). The air forced down into the soil was pushed out of six small holes made with a hot needle about 6 cm down from the lip of the pots. Before beginning the second measurement, the exhaust valve was turned to restrict air flow by 100% to force air down through the soil, and the flow was set to the maximum with desiccant on full scrub for one hour until the soil in the pot was fully dried. Pots were weighed before and after drying as the weight difference between wet and dry soil in the 65 mm pots was found to be approximately 30 g in a preliminary trial. A second, dry measurement, was then run in which all the above conditions were the same, except humidity and soil moisture were at 0%.

Another gas exchange experiment was conducted using a different set of seedlings. All settings were kept the same with the exception that seedlings were

acclimated and measured at a PAR of 300 μ mol m⁻² s⁻¹ at the plant level, and each measurement was only conducted for 24-hours on well-watered seedlings only. During this time the LI-COR 6400XT software malfunctioned, and a LI-COR 6800 with 6800-17 small plant chamber (Figure 1-2 in:

https://www.licor.com/documents/eg2bp3sqbr4a97wjwhvupgigqter9zic) was used instead following factory guidelines, with flow rate of 800 μmol s⁻¹, pressure valve set to 0.2 kPa, relative air humidity to 50%, sample CO₂ to 400 μmol mol⁻¹, fan speed of 10,000 rpm, and a block temperature of 20 °C. The auto log logged data was collected once every 300 seconds for 1440 minutes with matching after each log point. The LI-COR 6800 also uses a system for pushing air down through the soil, but instead of holes in the side of the pot, a silicon exhaust cup was attached to the bottom of the pot. This cup has a nipple that is opened or closed until the measured leak value of 20% is achieved.

Daughter cladode titratable acidity and isotopic mass spectrometric analysis

To determine daily nocturnal acidity build-up in daughter cladodes, 4year-old *O. ficus-indica* individuals in 11.3-1 pots within the University of Nevada, Reno greenhouse were selected for analysis. A 10 mm Fisherbrand Cork Borer was used to take punches of cladodes within different size categories ranging from 2-35 cm in length. In May of 2016 tissues were collected at the appropriate dusk and dawn times and immediately weighed and flash frozen for later titratable acidity analysis. Titratable acidity was measured and calculated exactly as described above in the titratable acidity of *O. ficus-indica* seedlings section.

A second set of samples was also taken in May of 2017 at noon from the same size categories using a 10 mm cork borer. These samples were weighed, and immediately lyophilized. The freeze-dried samples were then pulverized with a bead beater, and 2 to 3 mg of each sample was prepared for shipment to the FIRMS or SIF facility as described above for *O. ficus-indica* seedlings.

Daughter cladode gas exchange measurements

To measure the rate and duration of photosynthesis in growing daughter cladodes, emerging 2.5 cm cladodes were placed in a custom clear Plexiglass chamber (30cm tall and 17cm wide cylinder, Figure 1A; Tripp Plastics, Reno, NV) connected to the LI-COR 6400XT. The node of the daughter cladode was flush with the base of the chamber, and puddy (21-601, Ideal Industries) was used to seal any air gaps around the node. For this experiment the LI-COR mixer was set to hold the reference chamber CO₂ at 400 µmol mol⁻¹, the flow rate was set to 700 µmol s⁻¹, leaf fan to fast (setting 5), block temperature was set to 25 °C, and the ambient PAR was measured with the LI-COR 6400XT external light sensor placed in a level position above the chamber with puddy. At the beginning of each measurement, the length of the cladode was measured by looking across a measured grid on each side of the custom chamber, and then the LI-COR 6400XT was set to auto log ever 10 minutes for 1440 minutes while matching once ever 30 minutes. Measurements were repeated daily until the cladode grew too large to fit

in the chamber (about 15 cm in length). ΔCO_2 was calculated by subtracting the sample chamber CO_2 (PPM) from the reference chamber CO_2 concentration. All cladodes were measured when new cladodes were flushing between April 3rd and June 22nd in 2017.

In an attempt to separate out variation due to changing conditions within the greenhouse (standard daytime greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night with gradual sunup and sundown), 24-hour gas exchange of a growing daughter cladode on an *O. ficus-indica* individual grown under growth chamber conditions was also measured. The growth chamber (AR 75L, Percival Scientific, Derry, IA.). was set to the known optimum temperature of 25 °C/15 °C day/night temperature with a 12-hour day night cycle, and a measured canopy PAR level of 366 μ mol m⁻² s⁻¹.

Statistical Analysis

All plots and statistical analysis for this study were performed in PRISM 9 software (Graphcladode Software), except for plots and calculations for cladode length growth rate, which were performed using Excel software (Microsoft, Inc.).

Seedling titratable acidity samples were grouped by tissue type (cotyledon or cladode), samples taken at dawn *vs.* dusk, and age (30, 50, 75, and 100 days after germination) and plotted together in a grouped column graph. A three-way analysis of variance (ANOVA) was performed to determine if any significant variance was due to differences in tissue type, time of day, and age or any interactions among these variables. Welch's *t*-test was conducted to determine if significant differences were evident in nocturnal tissue acidity within each tissue type collected at dusk and dawn for each age group (eight independent comparisons). The sample size varied from five to ten within each group.

Seedling δ^{13} C isotope ratios were grouped by tissue type, and age (30, 50, 75, and 100 days after germination) and plotted together in a grouped column graph (n = 6). A two-way ANOVA was run to see if any significant variance in δ^{13} C ‰ vs. VPDB (Vienna Pee Dee Belemnite) occurred due to differences in tissue type and age, or an interaction between tissue type and age. Welch's *t*-test was used to determine if significant differences in δ^{13} C ‰ vs. VPDB occurred between tissue types within each age group (4 independent comparisons).

Daughter cladode tissue acidity was grouped into eight different size categories (0-2.5, 2.5-5, 5-10, 10-15, 15-20, 20-25, 25-30, and 30-35 cm) in cladode height with separate dawn and dusk tissue acidity measured in each category. Welch's *t*-tests were performed to determine if a statistical difference occurred between dawn and dusk acidity in each category (8 independent comparisons). The minimum sample size was 2 and max of 6.

Daughter C isotope ratios were grouped into 9 size categories of 0-2, 2-4, 4-6, 6-8, 10-12, 12-14, 14-16 cm and adult cladodes with an n=6. cladodes in the adult category were fully matured cladodes towards the base of the plant that were 35 cm or more in length. A one-way ANOVA was conducted to determine if any variance in measured δ^{13} C ‰ vs. VPDB occurred across the size categories. A follow up Tukey's multiple comparisons test was used to reveal any groupings among size categories in relation to measured δ^{13} C ‰ *vs.* VPDB.

Gas exchange results for seedlings grown in watered conditions under a PAR of 100 μ mol m⁻² s⁻¹, dry conditions under a PAR of 100 μ mol m⁻² s⁻¹, watered conditions under a PAR of 300 µmol m⁻² s⁻¹, and daughter cladodes in a custom transparent Plexiglas chamber were analyzed independently using the same analysis. All 24-hour measurements for each individual seedling or cladode were plotted together with y-axis being the photosynthesis rate (μ mol CO₂ m⁻² s⁻¹ for seedlings and ΔCO_2 (ppm) for daughter cladodes) and x-axis as time (h:m:s). All of the 24-hour measurements were aligned to the time point in which the attached light sensor recorded a PAR of 0 µmol m⁻² s⁻¹. Then, an area under the curve (AUC) analysis was performed for each of the 24-hour measurements and categorized by cladode length, which was the calculated total diel carbon assimilated for each cladode length measured (net μ mol CO₂ m⁻² for seedlings and net ppm CO_2 for daughter cladodes). Afterward, each plot was split into only light periods or dark periods (approximately 12 hours each). The AUC of the light period plots reflected the total carbon assimilated in the light, and the AUC of the dark period plots reflected the total carbon assimilated at night. Lastly, the calculated diel, light, and dark total carbon assimilated were all plotted together vs. measured cladode length for each measurement for each experiment to visualize trends in carbon fixation as *O. ficus-indica* cladodes grew.

To determine the growth rate of the central cladode in seedlings, the measured cladode length and combined cotyledon length was plotted *vs*. days

after germination. The rate of seedling cladode length increase was found to briefly decrease as the cotyledons began to senesce. So, the center cladode length growth rate was calculated by determining the slope of the plotted line before cotyledon senescence began, which was determined by photographs and the initial decrease and reduction in combined cotyledon length *vs*. days after germination. These values were averaged together for the PAR 100 seedlings (n = 4) and the PAR 300 seedlings (n = 4). These experiments were not directly comparable because PAR 100 seedlings had intermittent drying for the dry experiment, while PAR 300 seedlings were well watered for the duration of the experiment, but a standard *t*-test was used to measure for significant difference of average cladode length growth between the PAR 100 and PAR 300 experiments regardless.

To determine growth rate of daughter cladodes during the sample period, the length of each daughter cladode was plotted with the day it was sampled. The slope of the line was used to calculate cladode growth rate and averaged across all measured daughter cladodes (n = 5) except for the daughter cladode measured under growth chamber conditions.

Results

CAM development in seedlings

Seedling age, tissue type, and time of day all had a highly significant effect on titratable acidity (Table 1) with the interaction of days after germination with tissue type also having a significant effect on tissue acidity variance (Table 1). No significant difference in seedling tissue acidity of cotyledons collected at dusk *vs.* cotyledons collected at dawn was evident until 75 days after germination, and this difference was not significant again until 100 days after germination (Figure 2A). No significant difference in seedling tissue acidity of cladodes collected at dusk *vs.* cladodes collected at dawn until 75 days after germination was detected, and this significant difference persisted 100 days after germination (Figure 2A). Mean δ^{13} C ‰ ratio values for of cotyledons (-18.43 ‰) and cladodes (-17.53 ‰) were not significantly different 50 days after germination (Figure 2B). While these values were not indicative of strong CAM, they were well within the range of CAM species (O'Leary, 1988; Messerschmid et al., 2021)

In the watered PAR 100 gas exchange experiments, there was an increase in duration and rate of CO₂ fixation during phase 1 and 4, but not 2 and 3 as the central cladode grew (Figure 3A, Sup. Figures 1-6). Total net carbon assimilated in well-watered seedling grown under PAR 100 was negative during the dark and diel periods until the central cladode reached 5 cm in height (Figure 3B, Supplemental Figures 7-11). Net carbon assimilation in the light period was higher than the dark period (but still negative), until the central cladode was 4 cm in height, at which point net carbon fixation during the dark period surpassed the light period and increased steadily (Figure 3B). Seedlings grown under dry conditions under PAR 100 also showed an increase in CO₂ assimilation duration and rate during phases 1 and 4, but not phases 2 and 3 (Figure 4A, Supplemental Figures 12-15). However, the amount of CO₂ fixed in these phases was diminished compared to seedlings grown under well-watered conditions under PAR 100 (Figure 3A, B). The net diel CO₂ assimilation in seedlings grown under dry conditions under PAR 100 was negative until the cladodes reached 5 cm in height (Figure 4B, Supplemental Figures 17-20). The net nocturnal CO₂ assimilation was less than the net day CO₂ assimilation rate until the cladodes reached 5 cm in height, at which point the net CO₂ assimilated during the night surpassed the daytime fixation (Figure 4B).

Seedlings grown under well-watered conditions under PAR 300 showed a longer duration and higher rate of CO_2 assimilation in phases 1 and 4 (Figure 5A, Supplemental Figures 26-33) than that of seedling grown under well-watered conditions with PAR 100 (Figure 4A) and seedlings grown under dry under PAR 100 (Figure 3A). As observed for seedlings grown under PAR 100 under either well-watered or dry conditions, PAR 300 seedlings exhibited neither CO₂ assimilation in phase 2, nor obvious CO_2 assimilation during phase 3 (Figure 5A). Initially, most of the daily net CO₂ assimilated by seedlings under PAR 300 was fixed during the day in phase 4, but the nocturnal net CO₂ assimilated in phase 1 increased steadily (Figure 5B). A noticeable decrease in both daytime, and especially nighttime CO_2 assimilation was apparent as the cotyledons began to senesce from the seedlings under PAR 300 (Supplemental Figures 34-37). However, the occurrence of cotyledons senescing and falling off the seedlings was not frequent or consistent enough to make any clear-cut statements about this phenomenon. Unlike seedlings grown under PAR 300, seedlings grown under PAR 100 did experience cotyledon senescence during the sample period, but the cotyledons never fell off within the sampling time frame. The average cladode height growth rate was 0.101 (+/- 0.017) and 0.124 (+/- 0.027) cm/day for

seedlings grown under PAR 100 and PAR 300, respectively, but these rates were not significantly different (P value = 0.2078).

CAM development in daughter cladodes

Tissue samples taken from daughter cladodes revealed no significant difference in dusk/dawn tissue acidity until the cladodes grew to a height of 2.5-5 cm (Figure 6A). This difference in dusk/dawn tissue acidity increased as cladodes grew larger (Figure 6A). The average δ^{13} C ‰ ratio values for daughter cladodes ranged between -16.1 +/- 0.7 in adult cladodes, and -14.7 +/- 0.3 in cladodes 12-14 cm in height, which is typical of strong CAM (Figure 6B) (O'Leary, 1988). For daughter cladodes undergoing development with heights in size ranges of 0-2, 2-4, 4-6, 10-12, and 12-14 cm, the δ^{13} C ‰ ratio values (-17.08 to -14.02) were slightly less negative, but also typical of strong CAM (Figure 6B).

Initially, daughter cladodes showed no recognizable phases in CO₂ uptake over the 24-hour measurement period (Figure 7A). However, as the cladodes grew, the net diel, nocturnal, and daytime CO₂ assimilated first decreased, and then increased (Figure 7B). CAM phases 1, 2, and 4 periods of CO₂ assimilation in daughter cladodes increased as the cladodes grew larger in size (Figure 7A). Notably, no net positive diel CO₂ assimilation was detected until the daughter cladodes reached 7 cm in height or greater (Figure 7B). Both net nocturnal and daytime CO₂ assimilation increased as the daughter cladodes reached 15 cm (Figure 7A, B), although this trend was not seen in every cladode measured (Figure 7B, Supplemental Figures 42-46). Within the sample period, daughter cladodes showed a linear growth trend in relation to cladode height with an average growth of 1.00 cm (+/- 0.22) per day (Supplemental Figures 53-57). The daughter cladode measured on the plant grown under growth chamber conditions showed net positive diel and light period CO₂ assimilation, with net negative nocturnal assimilation at 8.5 cm in height (Supplemental figure 59), and both net light and dark period CO₂ fixation increased with the same magnitude as the cladode height increased. The growth chamber-grown daughter cladode showed clear periods of assimilation during phase 1 and 4, but only small, brief spikes of assimilation in phase 2 (Supplemental Figure 58). The measured cladode also had a linear gain in cladode height of 1.16 cm per day (Supplemental Figure 60).

Discussion

Unlike the tropical epiphyte *O. elatior* seedlings measured in Winter et al., 2011, *O. ficus-indica* seedlings showed no initial C₃ photosynthetic CO₂ assimilation pattern, which was active all day long during phase 3 when only cotyledons were present (Figure 3A, 4A, & 5A). Initial net diel CO₂ fixation was negative in seedlings grown under PAR 100 (Figure 2B & 2B), and initially positive in seedlings grown under PAR 300 (Figure 5B). While the rate and duration of CO₂ fixation during phase 1 and 3 increased as the seedlings grew, phase 2 CO₂ assimilation at dawn measured in *O. elatior* seedlings was not observed in any of the *O. ficus-indica* seedling gas exchange experiments. These differences are likely due to *O. ficus-indica* being a terrestrial species adapted to especially arid regions, while the epiphytic *O. eliator* is adapted to the canopy under more humid conditions.
While gas exchange measurements of *O. ficus-indica* seedling grown under dry conditions with PAR 100 did not mimic conditions of prolonged drought, they showed the non-acclimated response in quickly dried soil. While the seedlings fixed less overall CO₂ under dry experiment, CO₂ assimilation in phases 1 and 4 were still visible (Figure 4A). Net positive fixation of CO₂ was apparent as cladodes reached 5 cm in height (Figure 4B), and the majority of CO₂ was fixed at night (Figure 4A & B), demonstrating that photosynthesis continues for at least one day, even when the soil was completely dried. Repeating this experiment with prolonged drought and re-watering would be useful as *O. elatior* was shown to switch from mostly daytime CO₂ assimilation to nighttime assimilation when subjected to water-deficit stress for several days, and switch back when watered (Winter and Holtum, 2011). However, *O. ficus-indica* seedlings fixed CO₂ predominantly at night after 5 cm in length in all gas exchange experiments, suggesting that unlike *O. elatior*, *O. ficus-indica* predominantly uses CAM.

Number of days after germination, time of day, and tissue type all had a significant effect on measured tissue acidity in seedlings (Table 1). High tissue acidity of *O. ficus-indica* seedling cotyledons and cladodes at both dawn and dusk only 30-days after germination (Figure 2A) may have been a response to the gibberellic acid treatment (Swanson and Jones, 1996) used in the initial media germination recipe, although this has yet to be well demonstrated in many plant, let alone a cactus species. Like many constitutive CAM plant species, *O. ficus-indica* seedlings demonstrated an early onset of CAM documented through tissue acidity measurements (Altesor, 1993; Loza-Cornejo et al., 2003; Hernández-

González and Villarreal, 2007) as both the primary cladode and cotyledons had significantly higher tissue acidity at dawn then dusk 75 days after germination. However, this dawn/dusk difference was no longer significant in cotyledons at day 100 as they senesced, whereas the cladode dusk/dawn acidity remained significantly different as they continued to grow.

No difference in the timing, frequency (Supplemental Figure 61), or isotopic ¹²C: ¹³C ratio in seedling tissues was detected in O. ficus-indica grown on media containing sucrose vs. no sucrose. Collectively, O. ficus-indica seedlings displayed a mean germination rate of 63% using the methods outlined in this report. Other studies have also reported high germination rates when using nail clippers in combination with gibberellic acid treatment (Podda et al., 2017), and chemical scarification with concentrated H₂SO₄ or H₂O₂ for a few minutes (Altare et al., 2006; Khan, 2006). The latter method was found to be more challenging, as H₂SO₄ damaged many embryos within two minutes of application, and scarified germination rates were near zero in our hands. Interestingly, the fungal species Penicillium chrysogenum, Phoma medicaginis, and Trichoderma koningii, and *Trichoderma harzianum* have all been shown to promote germination in O. streptacantha, O. robusta, and O. leucotricha (Delgado-Sánchez et al., 2011; Delgado-Sánchez et al., 2013) and may be helpful for improving germination rates in O. ficus-indica as well.

A high occurrence of tricotyledonous seedlings, a three-cotyledon trait known to be caused by only a few mutated genes in *Arabidopsis thaliana*, was measured, but the molecular basis in *O. ficus-indica* is unknown (Khan, 2006).

Within our batch, we observed only one tricotyledonous seedling (Supplemental Figure 62B). Other O. ficus-indica seedling oddities were observed, such as several seedlings that had roots emerging from the point at which the central cladode connects with the hypocotyl (Supplemental Figure 62A), possibly for anchoring the central cladode to the ground when cotyledons are lost. We also observed two unfused seedlings germinate out of a single seed (Supplemental Figure 62C). Monozygotic polyembryony is common in angiosperms (Tisserat et al., 1979; Filonova et al., 2002). Two to three O. ficus-indica seedlings have been grown from single seeds in the past (Vélez-Gutierrez and Rodríguez-Garay, 1996), and gibberellic acid, used here, has been shown to promote polyembryony in O. ficus-indica (Kaaniche-Elloumi et al., 2013). We also observed one seedling with two fused cladodes connected to a single hypocotyl (Supplemental Figure 62D), and another that produced cotyledons that did not senesce, but failed to produce a central cladode. While these abnormal growth forms deserve further study and are a reflection of the genetic diversity of the particular seedlings used in this study, only intact, normal phenotypes were used for experimentation in this study.

Young, developing *O. ficus-indica* daughter cladodes had an increase in nocturnal tissue acidity build-up between 2.5 and 5.0 cm in length (Figure 6A), ¹²C: ¹³C ratios within the range of typical CAM regardless of size (Figure 6B), and an initial net negative dip in both light and dark fixation of CO₂ before emergence of a net positive assimilation in both the light and dark periods (Figure 7B) with the increase of carbon gain during phases 1, 2 and 4 (Figure 7A). Collectively, these results support the notion that as daughter cladodes emerge from mother cladodes, they initially start as sink tissues that metabolize resources partitioned from mature parts of the plant, releasing more CO₂ than fixed by either CA/PEPC or RUBISCO. These results are consistent with Wang et al., 1998 who found that daughter cladodes did not have a positive net CO₂ uptake until 20 days after initial appearance, that the sink-to-source transition is characterized by an increase in nocturnal malate build-up, that daughter cladodes imported carbohydrates from mother cladodes until 25 days after emergence, and was accompanied by a strong increase in sucrose synthase and hexokinase enzyme activities, which are linked to breakdown of carbohydrates from source tissues. The 24-hour gas exchange measurements in Wang et al., 1998 showed brief periods of phase 4 assimilation, and primarily phase 1 assimilation of CO₂ when the cladodes reached 28-days of age, while our results clearly showed increasing CO₂ assimilation in phase 1, 2, and 4 in growing daughter cladodes (Figure 7A).

While the greenhouse grown plants did not exhibit a net positive diel carbon assimilation until the cladodes reached about 10 cm in length (Figure 11B), this trend was not observed for all cladodes measured (Supplemental Figure 47-48). However, the daughter cladode measured on the *O. ficus-indica* individual grown in the growth chamber showed higher daytime than nighttime net CO₂ assimilation (Supplemental Figure 59), and both daytime and nighttime net CO₂ assimilation increased at a similar magnitude as the cladode grew, which was also observed by Winter et al., 2008 in lab conditions with similar PAR level. Unlike the greenhouse, the growth chamber daughter cladode had consistent light intensity during the light period and optimal day night temperatures. The resulting duration and rate of CO₂ fixation during phase 4 was much higher (Supplemental Figure 58) than that observed in greenhouse grown plants (Figure 7A, Supplemental Figures 42-46), which were likely limited in the amount of CO₂ that could be fixed during phase 3 and 4 by a gradual reduction of light at dusk, and gradual increase in light at dawn respectively. Curiously, while phase 4 was dramatically increased in the growth chamber grown plant, a strong increase in phase 2 *vs.* greenhouse plants was not observed (Supplemental Figure 58). More experimentation under growth chamber conditions is needed to make meaningful comparisons, but the greenhouse grown plants likely better represent cladode growth dynamics of plants grown under field conditions.

In this study, an early, significant build-up of nocturnal tissue acidity, typical CAM ¹²C: ¹³C ratios, and primarily nocturnal net CO₂ assimilation in *O*. *ficus-indica* seedlings all indicate the use of CAM in cotyledons and cladodes shortly after germination. In addition, *O. ficus-indica* daughter cladodes emerge as sink tissues that release more CO₂ through respiration than fixed by photosynthesis and begin to assimilate CO₂ both at night and briefly during dawn and dusk. These results show that seedlings use CAM photosynthesis early in development, and C₃ photosynthesis is not the predominantly used form of photosynthesis even under well-watered conditions. These findings also show that developing cladodes of *O. ficus-indica* begin as sink tissues, and switch to CAM after reaching about 5 cm in length, and fix CO₂ predominantly at night under greenhouse condition, but predominantly during the day under artificial conditions with a continuous light level throughout the day. C₃ photosynthesis does not appear to contribute to the high productivity observed in *O. ficus-indica* in developing tissues.

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Tables

Table 1: $\delta^{13}C$ ‰ ratio value ranges observed for different modes of

photosynthesis adapted from (O'Leary, 1988).

δ^{13} C ‰ values in C ₃ , C ₄ , and CAM plants.							
C ₃ Photosynthesis	C ₄ Photosynthesis	Typical CAM	Strong CAM	Wook CAM Plants			
Plants	Plants	Plants	Plants	weak CAIVI Flains			
-25 to -29	-12 to -16	-10 to -20	-11	-28			

Table 2: 3-way ANOVA results of Tissue acidity (μ mol H+ / g f.w.) of

cotyledons sampled at dusk and dawn, and cladodes at dusk and dawn at 30, 50,

75, and 100 days after germination.

Source of Variation	% of total	P value	P value	Significant?
	variation		summary	
Days after germination	10.87	< 0.0001	***	Yes
Cotyl vs. cladode	9.076	< 0.0001	***	Yes
Dusk vs. Dawn	15.84	< 0.0001	****	Yes
Days after germination x Cotyl vs.	3.636	0.0428	*	Yes
cladode				
Days after germination x Dusk vs.	2.46	0.1337	ns	No
Dawn				
Cotyl vs. cladode x Dusk vs. Dawn	1.018	0.1275	ns	No
Days after germination x Cotyl vs.	1.327	0.3848	ns	No
cladode x Dusk vs. Dawn				

Figures



Figure 1: *Opuntia ficus-indica* daughter cladode gas exchange and seedling germination. **A**: The custom-built gas exchange chamber for measuring 24-hour gas exchange of *Opuntia ficus-indica* daughter cladodes as they grow. **B**: A 2-year-old *O. ficus-indica* seed with the seed coat clipped enough that the embryo can make contact with germination media containing gibberellic acid, and so the embryo has enough room to leave the seed coat. **C**: Between seedling gas

exchange experiments, seedlings were kept in 65 mm LI-COR pots in a plastic container with water in the bottom. Notice the holes in the sides of the pots for root respiration air flow to be pushed out of during measurements with the LI-COR 6400XT whole *Arabidopsis* chamber exhaust restriction set up.



Figure 2: Seedling tissue acidity and δ^{13} C. **A:** Tissue acidity (µmol H⁺ / g f.w.) of cotyledons sampled at dusk (dark blue bars) and dawn (light blue bars); and cladodes at dusk (dark red bars) and dawn (light red bars) at 30, 50, 75, and 100 days after germination. The significance comparisons are the results of independent Welch's paired *t*-tests (p < 0.05) where 'ns' is not significant and '**' is p < 0.0021, where *n*=5-10. **B:** Measured ¹²C:¹³C isotope ratio (δ^{13} C ‰ *vs*. VPDB) of seedling cotyledons (blue bars) and central cladode (red bars) at 30, 50, 75, and 100 days after germination. Isotope ratio values for 30, 75, and 100 days (all shown as values of -1 δ^{13} C ‰ *vs*. VPDB) after germination are still being analyzed at UC Davis and are not included in this dissertation. The significance

comparisons are the results of independent Welch's paired *t*-tests (p < 0.05) where 'ns' is not significant, n = 6.



Figure 3: Well-watered PAR 100 gas exchange. **A:** 24-hour gas exchange measurements of *O. ficus-indica* seedling 3 (see Supplemental Figures 1-5 for all PAR 100 watered seedlings) grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in wellwatered conditions as the central cladode grows (dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² s⁻¹) was logged every 5 minutes. **B:** The combined diel (blue circles), light (yellow squares), and dark (black triangles) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in all seedlings grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 4: Dry PAR 100 seedling gas exchange A: 24-hour gas exchange measurements of *O. ficus-indica* seedling 5 (see Supplemental Figures 12-15 for all Dry PAR 100 seedlings) grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20° C in well-watered conditions as the central cladode grows (dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² s⁻¹) was logged every 5 minutes. Each measurement was taken after allowing the soil to completely dry out for one day. **B:** The combined diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻² s⁻¹ at 20 °C after drying out for one day.



Figure 5: PAR 300 seedling gas exchange. **A:** 24-hour gas exchange measurements of *O. ficus-indica* seedling 8 (see Supplemental Figures 26-33 for all PAR 300 seedlings) grown at a PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in wellwatered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² s⁻¹) was logged every 5 minutes. **B:** The combined diel (blue triangle), light (yellow circle), and dark (black square) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in all seedlings grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 6: Daughter cladode tissue acidity and δ^{13} C. **A:** Tissue acidity (µmol H⁺ / g f.w.) of *O. ficus-indica* daughter cladode tissues collected at dusk (blue bars) and dawn (orange bars). The significance comparisons are the results of independent Welch's paired *t*-tests (p < 0.05) where 'ns' is not significant, '*' is p < 0.0332, '**' is p < 0.0021, and '***' is p < 0.0002, error bars represent standard error (*n* = 2-6). **B:** Measured ¹²C:¹³C isotope ratio (δ^{13} C ‰ *vs.* VPDB) of seedling cotyledons (blue bars) and central cladode (red bars) at 30, 50, 75, and 100 days after germination. The letters indicate grouping of compared means by Tukey's test (p < 0.05), error bars represent standard error (*n* = 6). For both figures, samples were taken from 4-year-old plants grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 µmol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night



Figure 7: Daughter cladode gas exchange. **A:** 24-hour gas exchange measurements of *O. ficus-indica* daughter cladode 4 (see Supplemental Figures 42-46 for all daughter cladodes) in the custom-built chamber as the cladode grows (dark blue to light blue to light red to dark red) under standard greenhouse conditions with natural light at approx. 1,100-1,500 µmol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night. The Δ CO₂ (ppm) is equal to the sample chamber subtracted by the reference chamber and was logged every 10 minutes. **B:** Combined diel (blue circle), light (yellow square), and dark (black triangle) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladodes grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 µmol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.

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Chapter 3

The nitrogen preference of *Opuntia ficus-indica*: a sand culture snapshot.

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Abstract

Cactus pear (Opuntia-ficus indica (L.) Mill.) is an important agricultural crassulacean acid metabolism (CAM) species that has been gaining popularity as a source of food, forage, fodder, secondary products, and as a biofuel feedstock. The preferred source of nitrogen for this species, whether it be nitrate (NO_3) , ammonium (NH_4^+), or a combination of both, is not well understood. This chapter summarizes the investigation into the nitrate and ammonium preference of cladodes grown in sand culture with distilled water for one month and given a cross-factorial nutrient solution of 0.0, 2.5, 5.0, and 10.0 mmol of nitrate and/or ammonium for a month. Physiological measures were assessed including cladode growth, relative water content, chlorophyll, tissue acidity, soluble sugars, starch, nitrate, ammonium, glyoxylic acid, nitrate reductase activity, and nitrogen and carbon content. Relative steady-state transcript abundance of genes that encode enzymes involved in N metabolism and CAM were also determined. Significant differences were found in all measured variables except for cladode length, relative water content, and carbon content. cladodes provided with only distilled water produced no new cladodes, showed increased starch content, decreased soluble sugar content, and increased tissue acidity. Furthermore, these control

cladodes showed increased steady-state mRNA expression of the CAM-related genes encoding aluminum-activated malate transporter (ALMT) and phospho*enol*pyruvate carboxylase (PEPC) and higher steady-state mRNA expression of the nitrogen metabolism-related genes including glutamine oxoglutarate aminotransferase (Fd-GOGAT), glutamate dehydrogenase (NADH-GDH460), glutamine synthetase 2 (GS2), but lower expression of asparagine synthetase (AS) than nutrient supplied cladodes.

Introduction

Contemporary climate models predict that both the severity and duration of drought caused by anthropogenic climate change will increase worldwide by the mid to late 21st century (Naumann et al., 2018; Pokhrel et al., 2021). The resulting depletion of ground water stores is already having a direct effect on agricultural production, especially in the southwestern United States. With climate change, the habitable zones of many plant species are shifting (Feeley et al., 2020). Large agricultural crassulacean acid metabolism (CAM) species such as those within the *Agave* and *Opuntia* genus, are projected to have range expansions with climate change due to their many adaptations to xeric regions around the globe (Owen and Griffiths, 2014). *Opuntia-ficus indica (L.) Mill.* is well adapted to long periods of extreme drought due to stem succulence, extremely thick cuticle, root retention, and CAM photosynthesis (Davis et al., 2019). This species has a rich history of traditional uses for food, forage, fodder, and profitable secondary products (Stintzing and Carle, 2005). More recently, it's potential as biofuel feedstock, especially in the production of methane, has been recognized (Yang et al., 2015; do Nascimento Santos et al., 2016; Calabrò et al., 2018; Krümpel et al., 2020).

As an obligate CAM species, O. ficus-indica fixes most of its required carbon at night. CO_2 entering the cell through nocturnally open stomata (Males and Griffiths, 2017) is converted into bicarbonate by carbonic anhydrase followed by fixation into phospho*enol*pyruvate (PEP) into oxaloacetate (OAA) by a CAMspecific phosphoenolpyruvate carboxylase (PEPC) isozyme (Nimmo, 2000; Boxall et al., 2017). This PEPC is constitutively expressed with its malate inhibition relieved through phosphorylation by a dedicated, circadian controlled phosphoenolpyruvate carboxylase kinase (PPCK) (Hartwell et al., 1999; Taybi et al., 2000). The resulting OAA is converted malate by malate dehydrogenase and shuttled into the vacuole by a putative aluminum malate transporter (ALMT) (Borland et al., 2016; Pereira et al., 2017) for storage until sunlight is available. During the day, stored malate is released from the vacuole, and decarboxylated by NAD(P)-malic enzyme (Winter and Smith, 2021) or phosphoenolpyruvate carboxykinase (PEPCK). Storage of malate, and thereby CAM activity can be quantified by measuring the titratable acidity difference in tissue samples collected at dusk and dawn (Cushman et al., 2008). The released CO₂ is then fixed by the Calvin-Benson-Bassham cycle by ribulose-1,5-bisphosphate carboxylase/oxygenase (RUBISCO) as in the C₃ photosynthetic pathway.

One advantage of CAM photosynthesis is the high concentration of CO₂ within the mesophyll cells during the day thanks to malate decarboxylation

coupled with stomata closure (Osmond, 1978; Cockburn et al., 1979; von Caemmerer and Griffiths, 2009). In the presence of high intercellular CO_2 concentrations, RUBISCO fixes more CO_2 into 3-phosphoglycerate (3-PGA) while reducing the amount of O₂ fixed into 2-phosphoglycolate (2-PGA). 2-PGA is converted into glycoxylate, which must be shuttled between the chloroplast, peroxisome, and mitochondria to be metabolized in an energy costly process known photorespiration (Busch, 2020). CAM photosynthesis has been shown to reduce photorespiration in plants (Cushman and Bohnert, 1997; Lüttge et al., 2012), as photorespiration is often considered a wasteful process that limits productivity, although it is also considered to be a process that may balance ATP and NAD(P)H concentrations under excess energy conditions, and a process for shuttling carbon-containing compounds between vesicles for nitrogen assimilation (Busch, 2020). Another advantage of CAM is high water-use efficiency (WUE). CAM plants are often recognized as having a 6-fold greater WUE than that of C_3 photosynthesis plants, and a 4-fold greater WUE than that of C₄ photosynthetic species (Borland et al., 2009) thanks in part to reduced transpiration from nocturnal stomatal conductance, and many co-adapted traits that both further enhance CAM and WUE (Niechayev et al., 2019).

While nitrogen (N) metabolism in C₃ and C₄ photosynthesis species is well studied, a robust understanding of N metabolism in CAM plants is lacking (Pereira and Cushman, 2019). Plants can take up N in the form of inorganic nitrate (NO_3^-) and ammonium (NH_4^+) (Hachiya and Sakakibara, 2017), and in organic N forms such as urea ($CO(NH_2)_2$) and released biological matter, which is

often decomposed into ammonium by microbial communities in the soil (Habteselassie et al., 2013; Wan et al., 2014). Nitrate and ammonium interact with one another within the soil and can limit or enhance total N uptake in plants depending on the abundance and ratio of these two oppositely charged molecules (Hachiya and Sakakibara, 2017). In addition, soils containing too much nitrate or ammonium can alter plant cellular pH, which causes detrimental changes in basic cellular functions such as osmosis, diffusion, membrane stability, and enzyme activities (Feng et al., 2020). In most standard nutrient solutions, nitrate and ammonium are present in the millimolar range in a 1:1 ratio for ionic charge balance, or with more nitrate than ammonium (Smith et al., 1983; De Rijck and Schrevens, 1998) as there are many ammonium-sensitive plant species (Britto and Kronzucker, 2002). The optimal nitrate and ammonium concentrations for any given plant species is dictated by adaptation to a specific environment (Houlton et al., 2007; Wang and Macko, 2011). Dry, nitrate rich landscapes, plants tend to prefer nitrate, whereas wet, ammonium rich landscape, plants tend to prefer ammonium. Some species might adapt to changes in nitrate and ammonium availability in the soil within only a few generations, although the rate at which plants can adapt to new nitrogen sources appears to vary among domesticated crop species. (Daryanto et al., 2019), and was shown to be limited in wild African grass species (Wang and Macko, 2011).

In CAM plants, inorganic nitrate and ammonium can be assimilated into roots directly by enzymes, and then transported to mesophyll cells for fixation into amino acids (Pereira and Cushman, 2019). Root uptake is regulated by ammonium transporters (AMT) and nitrate transporters (NRT). Nitrate is reduced to nitrite (NO₂⁻) by nitrate reductase (NR) with NADH or ferredoxin (Fdx) reducing power in roots or shoots, respectively. Nitrite is highly oxidized and needs to be transported and/or reduced to ammonia (NH₃) by nitrite reductase (NiR) with Fdx in the stroma of shoot cells or NADH reducing power in the stroma of root cells. Glutamine synthetase (GS) combines ammonium with an acyl phosphate intermediate into glutamine in the chloroplast. Glutamine is then converted to 2-glutamate by glutamine oxoglutarate aminotransferase (GOGAT). Alternatively, ammonium can be converted to carbamoyl phosphate by carbamoyl phosphate synthetase (NAD-GDH) can convert NH₄⁺ directly into glutamate by combining with 2-oxogluterate and using NAD(P)H reducing power. Glutamine and ammonium can also be converted into asparagine by asparagine synthetase (AS) in ATP-rich regions of the cytosol.

Several studies have documented the productivity of *O. ficus-indica* under different fertilizer treatments in the field (Garcia de Cortazar and Nobel, 1992; Dubeux et al., 2006; Sánchez et al., 2012; Nkoi et al., 2021) and the commercial N input is between 50–100 kg ha⁻¹ year⁻¹ (Davis et al., 2019). However, these studies were conducted with a wide variety of N sources, soil types, and production goals (e.g., fruit, cladodes, seeds, and methane production); and did not reveal the nitrate and ammonium preferences of *O. ficus-indica*. Interestingly, a 4-fold increase in nocturnal acidity in *O. ficus-indica* was observed when chlorenchyma N% increased 3-fold, and higher chlorophyll content was observed in seedlings grown in concentrated Hoagland's solution over six months (Nobel, 1983). A nutrient index was developed to estimate productivity given different nutrient availabilities for cacti and agave species (Nobel et al., 1987), but this model is not species specific and does not specify between nitrate and ammonium for N input. To our knowledge, the only study that investigated nitrate *vs.* ammonium preference in *O. ficus-indica* was performed by (Vázquez et al., 2000). Through measuring nitrate and ammonium depletion from the sand, the authors found that plants initially took up more N when given ammonium than nitrate after 5, 10, and 15 days of treatment, but this difference was not significant after 20 days. Furthermore, plants accumulated significantly more biomass in the above ground tissue, and slightly more on average in root tissue when plants were given nitrate. The authors concluded that in hydroponic conditions, *O. ficus-indica* absorbed more N when supplied with nitrate than with ammonium, and when supplied with nitrate, the plants showed increased biomass production.

In this study, the response *O. ficus-indica* to varying amounts of nitrate *vs.* ammonium was investigated in a sand culture experiment. Mature cladodes were planted in pots containing sand, watered with distilled H₂O for a month, and then given one of 16 different nutrient treatments with modified Hoagland's nutrient solution that varied in nitrate and ammonium concentrations in a cross-factorial design, as well as a continued distilled H₂O treatment. After one month of applying nutrient treatments growth, relative water content, chlorophyll, tissue acidity, soluble sugars, starch, nitrate, ammonium, glyoxylic acid, nitrate reductase activity, nitrogen, carbon, and relative transcript abundance of genes

that code for enzymes involved in N metabolism and CAM were all measured in an attempt to gain an understanding of how *O. ficus-indica* responded to differences in nitrate and ammonium availability.

Methods

Greenhouse experimentation and sample collection

Prior to planting, 102 mature daughter cladodes were collected from 4-year-old O. ficus-indica plants located in the Valley Road Greenhouse Complex at the University of Reno, Nevada. The original mature plants were grown in threegallon pots containing a 3:1 ratio of Sunshine MVP soil mix (Sun GroHorticulture, Bellevue, WA, USA) and decomposed granite with the cladode placed about 5 cm into the soil. Plants were watered once or twice a week depending on season, treated monthly with both a Miracle Gro[®] fertilizer (Scott's MiracleGro, Inc., Marysville, OH, USA) as well as insecticide treatment (Marathon[®] 1% Granular, OHP, Mainland, PA, USA) every six months, and repotted on an annual basis. The collected daughter cladodes were allowed to callus for two weeks under greenhouse shaded conditions to prevent infection upon planting. cladodes were then planted in 11.3 L plastic pots containing a base layer of gravel and the remaining volume with sand (Sandbox Play Sand, Quikrete, Atlanta, GA) sterilized via autoclave set to a 40-minute dry cycle at 121°C and 15 PSI. All plants received 1L of distilled H₂O for 1 month prior to applying nutrient treatments to allow for acclimation and to leech any mineral nutrients out of the sand. After acclimation, 6 cladodes were randomly selected

for each nutrient treatment, and the position of each cladode within the greenhouse was also randomized to mitigate for any possible differences in greenhouse microclimate. Under standard greenhouse conditions, the natural light was approximately 1,100-1,500 μ mol m⁻² s⁻¹ and temperature was 28-32 °C day/17-18 °C night.

The experiment was conducted in a cross-factorial design with respect to nitrate and ammonium concentrations (Table5). Each treatment received 1L, twice a week, of an assigned modified full Hoagland's solution with 0.0, 2.5, 5.0, or 10.0 mMol of nitrate and/or ammonium that was adjusted to pH = 5.7-5.8 (Supplemental 1 for exact nutrient recipe of each treatment). In addition, a 17th treatment, which only received 1L of distilled H₂O twice a week, was included as a negative control. After one month of applying treatments, cladodes were collected to measure various parameters detailed below.

Growth measurements

cladode length, width, new cladode number, root length, and root number were all measured before beginning treatments (after the 1-month acclimation treatment), and after the one-month treatment period. Center cladode thickness was also measured with a digital caliper (IP54 caliper, Baleigh Industrial, Manitowoc, WI) after the treatment period.

Relative water content

A 2-cm diameter cork bore was used to collect tissue from each cladode for relative water content. Fresh samples were immediately weighed and submerged in distilled water for 24 hours. After soaking, tissues were weighed again to acquire the turgid weight. Lastly, samples were dried for 72 hours in a freeze drier (7755030, Labconco, Kansas City, MO) and weighed again to acquire the dry weight. The relative water content was calculated as:

RWC (%) =
$$\left(\frac{f-d}{t} - d\right) * 100(\%)$$

Where f is fresh weight, t is turgid weight, and d is dry weight all in grams. Chlorophyll content

Chlorophyll content was determined using a protocol modified from (Ni et al., 2009). 300 mg of frozen and ground tissue was placed into 15 ml Falcon tubes (430791, Corning, Corning, NY) and mixed with 5 ml of 80% acetone in the dark. Samples were then centrifuged at 3000 x g at 4° C for 15 min and preserving the supernatant. Samples were loaded into disposable cuvettes and the absorbance at 663nm for chlorophyll a (Ca) and 645 nm for chlorophyll b (Cb0 was measured using a Thermo Scientific nanodrop 2000 spectrophotometer. The chlorophyll content was calculated as:

$$Ca \ (mg/g \ sample) = (12.7 * A - 2.69 * B) * \frac{v}{1000} * w$$
$$Cb \ (mg/g \ sample) = (22.9 * B - 4.86 * A) * \frac{v}{1000} * w$$
$$Ca + b \ (mg/g \ sample) = (8.02 * A + 20.20 * B) * \frac{v}{1000} * w$$

Where A is absorbance at 663 nm, B is absorbance at 645 nm, v is volume of extract in ml, and w is weight of the sample in g.

Titratable acidity

To determine the nocturnal acid stored overnight within treatments, between 0.844-2.886 g fresh weight material was collected with a 2-cm diameter cork bore at dusk and dawn from each cladode with and flash frozen with liquid nitrogen. The titratable acidity was determined using a modified protocol (Gehrig et al., 2005). Collected tissue was ground in a mortar and pestle containing liquid nitrogen. 0.5 g of freeze-ground tissue from each sample was placed in pre-chilled 15 ml conical tubes. 10 ml of 50% methanol was added to each sample, and the top volume point marked with a marker. A small hole was made in the cap of each tube, and samples were boiled in an 80 °C water bath for 10 minutes. After boiling, samples were brought back to the marked level with distilled H₂O and centrifuged at 3,000 *x g* for 10 minutes. The decanted supernatant was then placed into 50 ml beakers and titrated to a pH of 7.0 for malate equivalent, and 8.4 for citrate equivalent with 10 mM KOH. The H⁺ equivalent at 7.0 and 8.4 was calculated by:

$$H^+$$
 equivalent($\mu mol \ H^+/gFW$) = $w\left(\frac{0.01}{v}\right) * 1000$

Where w is fresh weight of sample in grams, v is volume of 10 mM KOH added in ml.

For each sample, the total nocturnal buildup of malate and citrate was calculated by subtracting the dusk sample H⁺ equivalent from the dawn sample H⁺ equivalent.

Starch content and soluble sugars

The soluble sugars, glucose, fructose, and sucrose and non-soluble starch contents were analyzed exactly as specified (Gomez et al., 2007). To summarize, 10 mg of freeze ground tissue underwent a methanol extraction and chloroform phase separation. The top phase containing soluble sugars, and the lower phase containing starch were separated for appropriate analysis. The soluble sugar fraction was analyzed by conducting sequential enzyme assays that measure the production of NADH at 340 nm in a SpectraMax M5 multi-mode microplate reader (17000, SpectraMax, San Jose, CA) after the addition of glucose-6phosphate dehydrogenase (10165875001, Sigma-Aldrich, St. Louis, MO) for glucose content, hexokinase (11426362001, Sigma-Aldrich, St. Louis, MO) and phosphogluco isomerase (10128139001, Sigma-Aldrich, St. Louis, MO) for fructose content, and ß-fructosidase (14504, Sigma-Aldrich, St. Louis, MO) for sucrose content. The lower starch containing phase was hydrolyzed into glucose monomers by first autoclaving, and then the application of an amylglucosidase (11202332001, Sigma-Aldrich, St. Louis, MO) treatment. The freed glucose monomers were then determined by measuring the production of NADH at 340 nm after the addition of glucose-6-phosphate dehydrogenase (10165875001, Sigma-Aldrich, St. Louis, MO) as in the soluble sugar assay.
Nitrate reductase activity

NR activity assays, roots were collected from each cladode and the cortex was removed by hand before recording fresh weight. Prior to experimentation, a phosphate buffer was made by combining 500 ml of 0.1 m KH₂PO₄ with 400 ml of 0.1 M NaOH and adding more of 0.1 M NaOH until pH = 7.5 was achieved. 1 l of an incubation buffer was made by adding 970 ml of the phosphate buffer with 30 ml of *n*-propanol, and 100 mM KNO₃. The incubation buffer was heated in a water bath for 20 minutes at 30 °C and then placed in a 60 Sonic Dismembrator (F60, Fisher Scientific, San Diego, CA) and vacuum pump for 15 minutes to eliminate O₂ from the solution. The incubation solution void of O₂ was then kept in a water bath at 30 °C until fresh tissue was collected.

Approximately 0.5 g of *O. ficus-indica* cortex free root tissue was placed into 15 ml glass tubes. Six ml of the O₂-free incubation buffer was added to each sample and all samples were placed into a vacuum chamber for two 1-minute rounds to promote infiltration of tissues with incubation solution. All samples were kept in the dark or under foil for the remainder of the experiment to prevent nitrate degradation by light. 1 ml of incubation buffer from each sample was then pipetted into 2 ml microtubes to represent time point 0 (T0). T0 tubes were incubated at room temperature for one hour. The remaining samples were incubated for 1 hour in a 30 °C water bath, and 1 ml was transferred to a second set of tubes to represent the 60-minute time point (T60). 1 ml of O₂-free incubation buffer was added in a separate microtube as a blank for spectrophotometer readings and final calculation. In each microtube from the T0, T60, and blank, 30 μ l of 1% sulfanilamide in 3 M HCl was added and vortexed. Then, 300 μ l of 0.02% of N-(1-Naphthyl) ethylenediamine dihydrochloride in reagent water was added and vortexed, and samples were allowed to incubate for 30 min at room temperature. Lastly, samples were loaded into quartz cuvettes and measured at 540 nm in a Thermo Scientific nanodrop 2000 spectrophotometer using the appropriate buffer blank. A 345 mg NaNO₂ in 500 ml water solution was diluted to make 0, 1, 2, 4, 8, and 16 μ M NO₂-/l standard solutions.

The reaction rate of nitrate reductase in solution was calculated by first converting T0 and T60 to concentration to μ M using the equation of the best fit line of the standard absorbance readings. This calculated concentration was normalized by dividing it by the initial sample weight. Lastly, subtracting the normalized T0 concentration from the T60 concentration gave the μ M NO₂ produced per gram fresh weight of sample per hour by NR.

Nitrate content

Nitrate content was determined using a modified protocol (Cataldo et al., 1975). Briefly, 20 mg of freeze-dried tissue was resuspended in distilled water and incubated at 45 °C for 1 hour. Samples were then mixed and centrifuged at 5,000 x g for 15 min. 0.2 ml of supernatant was placed into a 50 ml flask with 5% salicylic acid in concentrated H₂SO₄ for 20 min at room temperature. 19 ml of 2 N NaOH was added to each sample to adjust pH \geq 12. Flasks were gently vortexed for 5 minutes and 100 µl of each sample was loaded into a 96-well clear polycarbonate, flatbottom microliter plate (3364, Corning, Corning, NY) and the absorbance was measured at 410 nm using a SpectraMax M5 multi-mode microplate reader (17000, SpectraMax, San Jose, CA). Samples were compared to a set of 8 standards containing between 0 and 60 mg of NO₃⁻ using a KNO₃⁻ standard solution and normalized by sample dry weight.

Ammonium content Glyoxylic acid content

Ammonium and glyoxylic acid content were quantified following (Bräutigam et al., 2007). About 50 mg of homogenized freeze-dried tissue was collected from each cladode. Samples were mixed with 1 ml of 100 mM HCl with 500 μ l of chloroform in 2 ml test tubes. Samples were then centrifuged at 12,000 *x g* for 5 minutes at 8 °C. The aqueous phase was transferred to a new set of test tubes containing 50 mg of acid-washed activated charcoal, mixed by elution, and centrifuged again at 20,000 *x g* for 5 minutes in 8 °C. 200 μ l of the charcoal washed supernatant went into the glyoxylate assay, and 200 μ l went into the ammonium assay workflows.

The glyoxylate samples were combined with 20 μ l of a 1% (v/v) solution of phenylhydrazine in 100 mM HCl and incubated at 95 °C in a hot water bath for 2 minutes and immediately cooled on ice for 6 minutes. 100 μ l of concentrated HCl was added to each sample. 225 μ l of each sample was loaded into a well of a 96-well clear flatbottom microliter plate, and the absorbance at 520 nm was measured exactly at 4, 5, and 6 minutes after the additions of 25 μ l of 1.6% K₃Fe(CN)₆ solution using a SpectraMax M5 multi-mode microplate reader. The 200 μ l of ammonium samples were diluted 1:1 with 100 mM HCl. 20 μ L of this solution was mixed with 100 μ L of 1% (w/v) phenol, 0.005% (w/v) sodium nitroprusside solution in water, and 100 μ L of 1% (v/v) sodium hypochlorite and 0.5% (w/v) sodium hydroxide. All samples were then incubated at 37 °C for 30 minutes, and the absorbance at 520 nm was measured in a SpectraMax M5 multi-mode microplate reader. Concentrations were calculated with the equation of a linear curve with 12 ammonium standards between 0- and 20-mM concentrations of ammonium sulfate.

Carbon and Nitrogen content

Total carbon and nitrogen content were determined by loading approximately 50 mg of ground freeze-dried tissue from each cladode into clay crucibles (2203-828, Leco, St. Joseph, MI) for elemental analysis with a Leco 928 combustion analyzer (Leco, St. Joseph, MI). Results were normalized on a weight basis and presented as the ratio of unit N per unit C (N:C Ratio).

RT-qPCR of CAM and nitrogen related genes

In order to measure the expression of CAM and nitrogen metabolismrelated genes within treatments, plant tissue was collected at noon with a 2 cm diameter cork bore from each cladode and then immediately frozen in liquid N₂ and ground to a fine powder using a mortar and pestle. 100 mg of ground frozen tissue was used for RNA extraction using a modified Qiagen RNeasy Plant Mini Kit (Qiagen Cat. No. 79254) protocol that included the addition DNase digestion, and Fruit-Mate (9192, Takara Bio Inc., Kusatsu, Shiga, Japan), a proprietary nonionic polymer that binds to polysaccharides and polyphenols. The addition of Fruit-Mate was necessary to perform RNA extractions on *O. ficus-indica* due to the naturally occurring high pectin content (Goycoolea and Cárdenas, 2003). RNeasy kit protocol was followed exactly as specified by the manufacturer with the addition of 1 ml of Fruit-Mate to the samples in step 2, and on-column DNase digestion using the RNase-free DNase kit as specified by Qiagen. The RNA concentration was measured with the Thermo Scientific nanodrop 2000 spectrophotometer, and potential RNA degradation during the extraction was checked by electrophoretic separation on a 1% agarose gel with Qiagen RNA sample loading dye (74904, Qiagen). cDNA of the extracted RNA transcripts was generated following iScriptTM Reverse Transcription Supermix for RT-qPCR protocol (1708840, Bio-Rad Laboratories, Hercules, CA).

For RT-qPCR analysis, primers were designed for *O. ficus-indica* genes related to CAM and nitrogen metabolism shown in Table 1. Real-time quantification was done following the SsoAdvanced Universal SYBR Green Supermix (172-5271, Bio-Rad Laboratories, Hercules, CA) protocol. The relative amounts of cDNA in each sample were determined on the basis of the threshold cycle (Ct) for each PCR product and normalized to both UBQ10 (Op_ fin19) and ACTIN7 (Op_ fin88560) Ct values (Pfaffl, 2001, 2004). Predicted localization of the final product of each gene was estimated by first translating the cDNA sequence to protein sequence using the Expasy translate tool (https://web.expasy.org/translate/). Then, the resulting protein sequence was analyzed using LOCALIZER software (http://localizer.csiro.au/) to generate a subcellular localization prediction (Sperschneider et al., 2017).

Statistical analysis

All raw data input and calculations described above were performed in Microsoft Excel. Analysis of variance for one factor One-Way ANOVA, Tukey test (a = 0.05), and boxplots with letters for significance were conducted in RStudio using datasets, ggplot2, multcompView, and dplyr packages. Graphcladode software was used to plot group bar graphs of soluble sugars, and chlorophyll content along with standard error.

Results

Growth measurements and relative water content.

No statistical differences were found in cladode length among the nitrate and ammonium treatments before and after the treatment period (Table 5), nor separate groupings among treatments (Figure 1). There was a significant difference in regard to cladode width among treatments (Table 5) both before treatment measurements, and the diH₂O treatment which had the lowest average widths, whereas the 5.0 + 2.5 (mMol Nitrate + Ammonium) and 10.0 + 5.0treatments having the highest average widths (Figure 2). A highly significant difference in cladode thickness among treatments (Table 5) was found with the lowest treatment averages being diH₂O and 0.0 + 0.0 (Figure 3). Treatments with the largest average cladode thickness were 5.0 + 2.5 and 2.5 + 10.0 (Figure 3). Difference in average number of new cladodes was highly significant among treatments (Table 5) and all nitrogen treatments had an average new cladode number above 0 (Figure 4) and 2.5 + 5.0, 0.0 + 10.0, and 2.5 + 10.0 all having the highest average new cladode number of 2, and before treatment samples, and diH₂O had no new cladodes. Primary root number was highly significantly different (Table 5) among treatments, and root number was also significantly different among treatments (Table 5). Treatments 0.0 + 0.0 and 0.0 + 10.0 had the highest average root numbers (Figure 5), whereas 10.0 + 0.0 had the highest average root length (Figure 6).

Relative water content averaged between 67.8 and 77.1% among treatments and no significant difference among treatments was found (Table 5, Figure 7).

Biochemical analyses

Chlorophyll

A slightly significant difference in Chlorophyll a + b content was observed among treatments (Table 6). However, no treatments were individually grouped by Tukey's test (Figure 8 and 9) with the maximum average combined in the 10.0 + 5.0 treatment, and the lowest in the 0.0 + 2.5 treatment. Chlorophyll a content showed a significant difference among treatments (Table 6), but all treatments were grouped together as well (Figure 8 and 10) with the maximum average content evident in the 10.0 + 5.0 treatment and the lowest in the 0.0 + 2.5treatment. Chlorophyll b content showed an extremely significant difference among treatments (Table 6) with the highest average content in the 0.0 + 10.0 treatment, and lowest in the 0.0 + 5.0 treatment (Figure 8 and 11).

Titratable acidity

Dawn-dusk delta H⁺ to pH 7.0 (μ mol/gFW) showed a significant difference among treatments (Table 6) with the highest tissue acidity measured in the diH₂O treatment (Figure 12). Dawn-dusk delta H⁺ from pH 7.0 to pH 8.4 (μ mol/gFW) also showed a significant difference among treatments (Table 6), with the highest average tissue acidity evident in the 5.0 + 5.0 treatment (Figure 12).

Starch and sugar

Starch content among treatments showed a significant difference (Table 6), with the highest measured content in the diH₂O treatment (Figure 13). All treatments showed consistently higher average fructose than glucose, and glucose than sucrose (Figure 15, 16, 17, & 18). Individually, all three soluble sugars were statistically different among treatments (Table 5), with the highest concentrations in the 0.0 + 2.5 treatment, and the lowest in the diH₂O treatment.

Nitrate reductase activity

Preliminary results demonstrated that *O. ficus-indica* roots have a higher NR activity (average of 171.0 nmoles NO₂/g fw/h) than cladode chlorenchyma (21 nmoles NO₂/g fw/h) and hydrenchyma (16 nmoles NO₂/g fw/h) tissues. Roots

with cortex tissues were also shown to have less NR activity than that of roots with the cortex physically removed before analysis. Nitrate reductase (NR) activity measured in the roots of the plants submitted to different treatments was significantly different (Table 6). The highest NR activity was measured in the 10.0 + 2.5 treatment, while the lowest was observed in the 2.5 + 5.0 and diH₂O treatments (Figure 19).

Nitrate, ammonium, and glyoxylic acid

Nitrate and ammonium content showed significant differences among treatments (Table 6), with the highest recorded average nitrate content in the 10.0 + 0.0 treatment, and lowest in the 0.0 + 2.5 treatment (Figure 20). On the other hand, the 5.0 + 10.0 treatment showed the highest average ammonium content, while the 2.5 + 0.0 treatment presented the lowest (Figure 21). Glyoxylic acid content was significantly different among treatments (Table 6). The diH₂O treatment had the highest average glyoxylic acid content (Figure 22), but with a wide sample variance (2.0 to 5.5 µmol/mg sample).

Nitrogen and carbon

Nitrogen:Carbon ratio (N:C) among treatments was significantly different (Table 6). Interestingly, while the percent C was not significantly different (Table 6, Figure 24), the percent N (Figure 25) showed a significant difference (Table 6) with the highest percent N in the 10.0 + 2.5 treatment and N:C ratio occurring in the 5.0 + 10.0 treatment.

mRNA abundance of CAM and nitrogen metabolism-related genes

All measured relative expression levels for each gene were significantly different among treatments (Table 7, Figure 26). Of the CAM enzymes surveyed, the steady-state transcript abundance of ALMT and PPC1 were the highest in the diH₂O treatment (Figure 26, 27 and 28). In contrast, the relative steady-state transcript abundance of PPCK was highest in the 0.0 + 2.5 and 2.5 + 5.0 treatments, and lowest in the 0.0 + 0.0 treatment (Figure 26 & 29).

For N metabolism genes relative steady-state transcript abundance for NR was the significantly increased in the 10.0 + 0.0 and 10.0 + 2.5 treatments and the lowest in the 0.0 + 10.0 treatment (Figure 26 & 30). Nitrite Reductase (NiR) steady-state transcript abundance was highest in the 10.0 + 10.0 treatment, and lowest in 0.0 + 10.0 treatment (Figure 26 & 31). GOGAT steady-state transcript abundance was significantly increased in the 10.0 + 10.0 and diH₂O treatments, and lowest in the 5.0 + 10.0 treatment (Figure 26 & 32). AS steady-state transcript abundance was highest in the 0.0 + 5.0 treatment and lowest in the 5.0 + 10.0 treatment (Figure 26 & 32). AS steady-state transcript abundance was highest in the 0.0 + 5.0 treatment and lowest in the 5.0 + 10.0 treatment (Figure 26 & 33). GDH460 steady-state transcript abundance was highest in the 2.5 + 5.0 treatment, and lowest in 5.0 + 0.0 treatment (Figure 26 & 34). GDH201910 steady-state transcript abundance was highest in the diH₂O (Figure 26 & 35). Lastly, GS30900 showed no significant difference steady-state transcript abundance among the treatments (Figure 26 & 36). In contrast, GS94700 had about a 3-fold higher steady-state

transcript abundance in di H_2O than that of all of the other treatments (Figure 26 & 37).

Discussion

All treatments put on new cladodes with the exception of diH₂O treatment controls (Figure 4). The formation of new biomass might have affected the measured concentration of nitrate and ammonium due to mobilization of nutrients between source tissue in mother cladodes and sink tissue in daughter cladodes (Marschnert et al., 1997). In O. ficus-indica chlorophyll content has been shown to increase with increasing amounts of nitrate (Nerd and Nobel, 1995), and to decrease under high light and elevated CO₂ conditions (Cui and Nobel, 1994). The differences observed in combined chlorophyll content among treatments was slightly significant (Table 6, Figure 8 and 9), with the difference in chlorophyll a (Figure 8 & 10) and b (Figure 8 & 11) being more significant among treatments. In most plants, including O. ficus-indica, limited N in the soil promotes root growth (Vázquez et al., 2000; Kiba and Krapp, 2016). An increase in average root length (Figure 6) with less N was not obvious, but the primary root number was higher in the 0.0 + 0.0 treatment (Figure 5). cladode width (Figure 2) and cladode length (Figure 1) did not vary greatly among treatments, but cladode thickness did (Figure 3). Curiously, cladode thickness has been shown to correlate strongly with relative water content (Scalisi et al., 2016), but relative water content did not vary among treatments within this experiment (Figure 7).

In O. ficus-indica, fructose, glucose, and sucrose levels make up 35%, 32%, and 33% of the relative sugar content, respectively, found in chlorenchyma, and 44%, 43%, and 13%, respectively, in the parenchyma under well-watered conditions (Nerd and Nobel, 1991). Homogenized samples (combined chlorenchyma and parenchyma) were analyzed for soluble sugar analysis. We found that average fructose content was higher than glucose content in all nitrate vs. ammonium treatments (Figure 15), and sucrose content was minimal (≤ 2.5 mg/mg sample) in all treatments (Figure 18). Interestingly, no measured soluble sugar content was observed in the diH₂O control treatment samples (Figures 15,16,17, and 18). The control treatment samples also had the highest starch content (Figure 14). These observations suggest that under nutrient limiting conditions, O. ficus-indica is likely storing soluble sugars by conversion into starch until nutrient availability becomes more favorable, as has been seen in other plant species (Christopher and Holtum, 1996; Rosa et al., 2009; Tao et al., 2013; Tao et al., 2017)

Vázquez *et al.* (2000) observed a significantly higher uptake of N from sand in *O. ficus-indica* given nitrate *vs.* ammonium until 20 days after application, at which time the authors suggested that a significant difference was not seen because most of the supplied N had been taken up from the sand. The authors also measured higher above and below ground biomass when *O. ficus-indica* was provided with nitrate *vs.* ammonium. Our results complement these former results in demonstrating that percentage N appears to be slightly higher in *O. ficus-indica* 2.5 ± 0.0 vs. 0.0 ± 2.5 , and 5.0 ± 0.0 vs. 0.0 ± 5.0 treatments. However, no statistically significant differences were observed in the 10.0 ± 0.0 vs. 0.0 ± 10.0 treatments (Figure 25). In addition, *O. ficus-indica* results showed a higher percentage of N when supplied with both nitrate and ammonium except in the 10.0 ± 5.0 treatment (Figure 25), which might be a result of nitrate toxicity (Islam et al., 2020). Nitrate and ammonium content in cladodes were both significantly different across treatments (Figure 20 and 21), but understanding these differences is complicated by the fact that ammonium and nitrate are readily interconvertible in both the roots and cladodes (Krapp, 2015; Taiz et al., 2015).

Prior to this study, nitrate reductase activity and nitrate content had been measured in *O. ficus-indica* cladodes and roots under both field and glasshouse conditions (Nerd and Nobel, 1995). Both Nerd and Nobel, 1995 and this study demonstrated increased nitrate content in cladodes (Figure20), and NR activity in roots (Figure 19) when given nitrate concentrations were increased. However, our study revealed that NR activity did not always increase when both nitrate and ammonium were present, specifically in the 2.5 + 2.5, 2.5 + 5.0, and 2.5 + 10.0 treatments (Figure 19). NR activity was observed under higher nitrate concentration, which is likely explained by the fact that NO₃⁻ is the substrate for NR. Nerd and Nobel, 1995 also found the highest NR activity in new cladodes *vs*. that in basal cladodes, and roots, which had the least amount of NR activity. Conversely, in preliminary measurements, we found that root tissue had the highest amount of NR activity compared with both chlorenchyma and parenchyma cladode tissues (data not

shown). This difference is likely because cortical root tissue was removed before conducting NR activity assays, and Nerd and Nobel 1995 used intact roots for measurements. NR in the roots reduces nitrate to nitrite using ferredoxin reducing power prior to transport to photosynthetic tissues, while the nitrate reductase in photosynthetic tissues uses NADH reducing power (Pereira and Cushman, 2019). The NR activity measured in this study likely represents the conversion of nitrate to nitrite before being converted into ammonium by nitrate reductase prior to being assimilated into amino acids or transported to the shoots, while NR activity in cladodes is likely likely to the conversion of nitrate to nitrite just prior to assimilation into amino acids via the GOGAT cycle in plastids (Krapp, 2015). Thus, in *O. ficus-indica*, both root and cladode NR activities increased with an increase in supplied nitrate (Figure 19) (Nerd and Nobel, 1995).

As in facultative CAM plants, which have the ability to switch from C₃ photosynthesis to CAM photosynthesis under unfavorable conditions, a switch to CAM has been demonstrated in (Winter et al., 2015) when either high concentrations of an unfavorable N source is present, and/or when a favorable source of N is limited (Pereira and Cushman, 2019). As an obligate CAM plant, *O. ficus-indica* is expected to increase CO₂ uptake by CAM as has been shown in other cactaceae species (Nobel, 1983). Indeed, steady-state transcript abundance for PEPC1 increased when ammonium increased in the 0.0 + 2.5, 0.0 + 5.0, 0.0 + 10.0 treatments and when nitrate increased in the 2.5 + 0.0, 5.0 + 0.0, 10.0 + 0.0 treatments (Figure 28), although this trend was not apparent in the combined nitrate and ammonium treatments. ALMT steady-state transcript abundance was

also higher in 0.0 + 5.0 and 10.0 + 10.0, (Figure 27). An increase in ALMT with increases in N might translate to an increase in transport of malate in and out of tonoplasts within these treatments, as metabolism in general may be increased in this situation (Kovermann et al., 2007). No obvious trends were recognized in PPCK steady-state transcript abundance among treatments in relation to nitrate and ammonium concentrations (Figure 29) even though there were significant differences among treatments for PPCK steady-state transcript abundance. Interestingly, diH₂O control plants showed the highest accumulation of organic acids (malate + citrate) (Figure 12), and relative expression of ALMT (Figure 27) and PPC1 genes (Figure 29), which suggests that CAM increased without nutrient supplementation. However, organic acid build-up in the diH₂O treatment might also be due, in part, to a stress response (Lopez-Bucio et al., 2000), rather than an increase in net CO₂ fixation, which could be revealed by diel gas-exchange measurements (Niechayev et al., 2019). Glyoxylic acid content (Figure 22) was the same across all treatments except for the diH₂O control supporting the possibility that photorespiration rates in O. ficus-indica remained similar regardless of N supply.

NR and NiR from homogenized cladode tissue showed a similar relative expression pattern among treatments (Figures 30 & 31), which was likely because a coordination of both enzymes for the initial fixation of nitrate is required (Sestili et al., 2018). In addition, the steady-state mRNA abundance of these two enzymes has been shown to increase when N supply is increased, especially in the form of nitrate. In the next step of the pathway, GS fixes ammonium into glutamine both

in the cytoplasm through GS1 (GS30900), and chloroplast with GS2 (GS94700) (Bernard and Habash, 2009). The expression of GS30900 and GS94700 were both relatively similar across treatments with the exception that GS94700 showed increased relative steady-state transcript abundance in the diH₂O control treatment (Figure 37), while GS30900 relative expression was similar in all treatments (Figure 38). This observation suggests that chloroplast GS is upregulated more so than cytosolic GS under nutrient deprivation in O. ficus-indica. Likewise, there was also an increase in GOGAT steady-state transcript abundance within the diH₂O control treatment (Figure 32). GOGAT mRNA expression was also significantly higher in the 10.0 + 10.0 treatment (Figure 32), suggesting that GOGAT expression in O. ficus-indica was higher when the most N was supplied, and when nutrient limited. The high relative mRNA expression of GOGAT in the 10.0 + 10.0 treatment was likely due to an increased fixation of glutamine into glutamate with more nitrogen availability, while the high GOGAT relative mRNA expression in the diH₂O control treatment might be due to glutamate production. Glutamate production has being linked to maintenance of redox homeostasis and ATP production via glycolysis when malate levels are high or NAD-MDH function is lacking (Selinski and Scheibe, 2019), as high malate levels (H⁺ equivalent) were also observed in the diH₂O treatment (Figure 12). Asparagine synthetase steady-state transcript abundance in the cytosol was highest in the 0.0 + 5.0 treatment, but interestingly lower in the high N treatments 10.0 + 10.0, 2.5 +10.0, 5.0 + 10.0, and the diH₂O control treatment (Figure 33). Under high concentrations of ammonium, GDH converts ammonium into glutamate

(Skopelitis et al., 2006). In this experiment, the highest steady-state transcript abundance of GDH460 was measured in the 2.5 + 5.0 treatment (Figure 34), and the highest GDH201910 steady-state transcript abundance was observed in the 2.5 + 10.0 treatment (Figure 35). GDH460 showed higher steady-state transcript abundance in the diH₂O control treatment compared to GDH20190, and both of these genes showed slightly higher steady-state transcript abundance in the 0.0 +5.0 treatment than the 0+10 treatments, corroborating previous results observed by (Skopelitis et al., 2006).

After one month of acclimating O. ficus-indica cladodes to sand culture with diH₂O and one additional month of varying nitrate and ammonium concentrations in applied nutrients solutions, significant differences were found among treatments for all independent variables measured with the exception of cladode length, relative water content, and percent carbon (Table 5, 6, & 7). Thus, the one-month acclimation and one-month treatment periods were long enough to elicit significant differences in growth, biochemical parameters, and gene expression in O. ficus-indica. Obvious trends and best overall nitrate vs. ammonium treatments were not clearly revealed for all the parameters studied perhaps because only a single, terminal measurement was taken. The experimentation described here might have been more informative if several tissue samples had been taken over the 2-month period to reveal a continuum of changes within each treatment. If multiple samples are taken more O. ficus-indica individuals may be needed to prevent variation due to destructive sampling for tissue samples. Despite this, these results showed that nutrient limited in O. ficus*indica* failed to add new cladodes, exhibited increased starch content, near complete reductions in the contents of several soluble sugar, and increased steadystate transcript abundance for ALMT, PPC1, GOGAT, GDH460, and GS genes. Overall, these results demonstrate that fertilizers designed for *O. ficus-indica* production should have either equal parts nitrate and ammonium, or slightly more nitrate than ammonium based upon the results obtained here and suggested by previous literature reports.

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Tables

Table 1: Recipes for nutrient working solutions that were stored in the dark at

 4°C and combined to make each treatment nutrient mix prior to application.

Reagent (pH	Molecular	Stock	g/500mL	Working	Working	Working	Micronutrients (pH 5 7-5 8)	g/500ml	Fe-EDTA (pH 5 7-5 8)	g/250mL
5.7-5.8)	Weight	Solution	(Stock	Solution	Solution	Solution	(pri 5.7-5.6)		(pii 5.7-5.6)	
			Solution)	(mL/1.5L)	(mL/1.5L)	(mL/1.5L)				
MgSO ₄ +	246.47	500mM	61.62	6.0	6.0	6.0	H ₃ BO ₃	1.43	EDTA	6.5
7H ₂ 0				(2.0mM)	(2.0mM)	(2.0mM)				
KNO3	101.1	500mM	25.28	2.52	5.01	10.02	$MnCl_2 + 4H_2O$	0.91	FeSO ₄ +	6.25
				(0.84mM)	(1.67mM)	(3.34mM)			/1120	
Ca(NO ₃) ₂ +	236.15	500mM	59.04	2.49	4.98	9.99	$ZnSO_4 + 7H_2O$	0.11		
4H ₂ O				(0.83mM)	(1.66mM)	(3.33mM)				
K_2SO_4	174.26	500mM	43.57	3.75	2.52	5.01	CuSO ₄ + 5H ₂ O	0.04		
				(1.25mM)	(0.84mM)	(1.67mM)				
(NH ₄) ₂ H ₂ PO4	132.06	500mM	33.02	3.75	7.5	15	H ₂ MoO ₄	0.01		
				(1.25mM)	(2.5mM)	(5.0mM)				
CaCl ₂	110.98	500mM	27.75	7.5	5.01	9.99				
				(2.5mM)	(1.67mM)	(3.33mM)				
NaH ₂ PO ₄	120	500mM	30	11.25	7.5	15				
				(3.75)	(2.5mM)	(5.0mM)				

Table 2: Amount of working solutions combined to make each treatment in ml.

The resulting nutrient solution was brought to a final volume of 1.5L with

distilled water. 250 ml of the final solution was added to each of the six OFI

individuals in each treatment twice a week for one month after.

	$0.0 \mathrm{mM}$	2.5mM	2.5mM	10.0mM	10.0mM	0.0mM	$0.0 \mathrm{mM}$	$0.0 \mathrm{mM}$	2.5mM	2.5mM	2.5mM	10.0mM	5.0mM	5.0mM	5.0mM	10.0mM	diH ₂ O
	NO3 ⁻ +	NO3 ⁻ +	NO3 ⁻ +	NO3 ⁻ +	NO3 ⁺ +	NO3 ⁻ +	NO3 ⁻ +	NO3 ⁻ +	NO3 ⁻ +	NO3" +	NO3 ⁺ +	NO3 ⁻ +	NO3 ⁻ +	NO3 ⁻ +	NO3 ⁺ +	NO3 ⁻ +	
	0.0mM	0.0mM	0.0mM	0.0mM	10.0mM	2.5mM	5.0mM	10.0mM	2.5mM	5.0mM	10.0mM	2.5mM	2.5mM	5.0mM	10.0mM	5.0mM	
	NH_4^+	NH_4^+	$\mathrm{NH_{4}^{+}}$	\mathbf{NH}_4^+	\mathbf{NH}_4^+	NH_4^+	NH_4^+	NH_{4}^{+}	NH_4^+	NH_4^+	\mathbf{NH}_4^+	$\mathrm{NH_{4}^{+}}$	NH_4^+	NH_4^+	$\mathrm{NH}_{4^{+}}$	$\mathrm{NH_{4}^{+}}$	
MgSO ₄	6	6	6	6	6	6	5	6	6	6	6	6	6	6	6	6	0
KNO3	0	2.52	5.01	10.02	10.02	0	0	0	2.52	2.52	2.5	10.02	5.01	5.01	5.01	10.02	0
Ca(NO ₃) ₂	0	2.49	4.98	9.99	9.99	0	0	0	2.49	2.49	2.49	9.99	4.98	4.98	4.98	9.99	0
K_2SO_4	5.01	3.75	2.52	0	0	5.01	5.01	5.01	3.75	3.75	3.75	0	2.52	2.52	2.52	0	0
CaCl ₂	9.99	7.5	5.01	0	0	9.99	9.99	9.99	7.5	7.5	7.5	0	5.01	5.01	5.01	0	0
NaH ₂ PO ₄	15	15	15	15	0	11.25	7.5	0	11.25	7.5	0	11.25	11.25	7.5	0	7.5	0
(NH ₄) ₂ H ₂ PO4	0	0	0	0	15	7.5	7.5	0	3.75	7.5	15	3.75	3.75	7.5	15	7.5	0
Micronutrients	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	0
Fe-EDTA	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	1.5	0

Table 3: The factorial design to test the response of OFI to understand possible synergistic effects with differences in NO_3^- and NH_4^+ availability. An additional treatment that only received deionized water was included (diH₂O).

Treatments	0.0mM NO₃ [.]	2.5mM NO₃ ⁻	5.0mM NO₃ [.]	10.0mM NO ₃ -
0.0mM NH₄+	0.0mM NH4 ⁺	0.0mM NH ₄ +-	0.0mM NH₄⁺-	0.0mM NH ₄ +-
	0.0mM NO3 ⁻	2.5mM NO ₃ -	5.0mM NO₃ ⁻	10.0mM NO ₃ -
2.5mM NH₄⁺	2.5mM NH ₄ +-	2.5mM NH ₄ +-	2.5mM NH₄⁺-	2.5mM NH₄⁺-
	0.0mM NO ₃ -	2.5mM NO ₃ -	5.0mM NO₃ ⁻	10.0mM NO₃⁻
5.0mM NH₄⁺	5.0mM NH ₄ +-	5.0mM NH4 ⁺ -	5.0mM NH4+-	5.0mM NH ₄ +-
	0.0mM NO ₃ -	2.5mM NO3 ⁻	5.0mM NO3 ⁻	10.0mM NO ₃ -
10.0mM NH₄⁺	10.0mM NH ₄ +-	10.0mM NH ₄ ⁺ -	10.0mM NH ₄ +-	10.0mM NH4 ⁺ -
	0.0mM NO ₃ -	2.5mM NO ₃ ⁻	5.0mM NO ₃ -	10.0mM NO3 ⁻

Table 4: List of primers designed for RT-qPCR analysis with predicted

subcellular localizations.

Gene	Primer	Description	Localized to	Primer Sequence	Annealing Temperature (°C)	Amplicon Lenth (bp)
Op_fin206820	ALMT10	Aluminum Activated	Mitochandria, Tonoplast	F:TTTCCTATCTGGGCTGGC	56.2	68
		Malate Transporter		R:TTTCAAGGGTGTTGCTGCTTC		
Op_fin7190	PPC1	Phosphoenolpyruvate	Cytosol	F:AGGCACAATGATGTGATGGA	55.2	150
		Carboxylase		R:CCAATCTCCCCAAAACTGAAGAA		
Op_fin211860	PPCK1	Phosphoenolpyruvate	Cytosol	F:AACCAGTGTTGACGAGCCA	60	154
		Carboxylase Kinase		R:ATGCCATCTACCGTACCAGC		
Op_fin52570	NR	Nitrate Reductase	Cytosol	F:CCACACTTTGACCCTGTCC	56.2	87
				R:TCCTCTGTCCGATTGGCGTA		
Op_fin241390	NiR	Nitrite Reductase	Chloroplast	F:ACAATAGGCACCAAGTCC	53.6	151
				R:GCAACCCATGAAACCGATAT		
Op_fin81140	GOGAT	Glutamate Synthase	Chloroplast	F:GTAAAGTTGAACCTGCTTCATCTA	53.6	124
				R:GAACAGATTGGGTGGAAAATC		
Op_fin236590	AS	Asparagine Synthase	Cytosol	F:ATCCCACTCCGTCACTGAAC	56.2	88
				R:ATCATCATCAAAGGCATTGCG		
Op_fin460	GDH	Glutamate Dehydrogenase	nase Cytosol	F:AGTGGGTTAGGGATGAATT	60	105
				R:AAGAAATTGACTTGGAAATTGGA		
Op_fin201910	GDH	Glutamate Dehydrogenase	Cytosol	F:ACCAAGAGATCCACCAAGG	60	123
				R:TTTGTCCCCATATCAGGTGC		
Op_fin30900	GS	Glutamine Synthetase	Chloroplast	F:GGAATGAAAGAAGGTTGACTGGA	55.2	82
				R:GTCGCTAATCGTGGTTGCTC		
Op_fin94700	GS	Glutamine Synthetase	Cytosol	F:AATCACCTCGTAGCCACC	55.2	121
				R:AAGCACCACTCCAGCAATC		
Op_fin19060	UBQ10	Ubiquitin	Mitochandria, Cytosol	F:CTTCTGGATGTTGTAGTCAGC	56.2	150
				R:GCTCTCAACCTCCAAAGT		
Op_fin88560	ACTIN7	Actin	Cytosol	F:CACATCTGTTGGAAGGTGC	53.6	139
				R:ATTTCTTTGCTCATACGGTCAG		

Table 5: Ordinary one-way analysis (ANOVA) results of growth and relative

water content. Significance codes: extremely significant 0 '***', highly significant

p<0.001 '**', sig	nificant p<0.01 '*'	, slightly significant	p<0.05 '.'.,	and not
significant p>0.	05 'NS'.			

Independent Variable	N (# within treatment)	DF (Treatment number)	F-value	P-Value	Significance code
cladode Length	3	17	1.569	0.1260	NS
cladode Width	3	17	2.074	0.0326	*
cladode Thickness	6	16	2.678	0.001	**
New cladode #	3	17	2.689	0.0063	**
Primary Root Length	3	17	2.544	0.0091	**
Root Length	3	17	2.097	0.0306	*
Relative water Content	6	16	1.533	0.1070	NS

Table 6: Ordinary one-way analysis (ANOVA) results of biochemistry measurements with 6 samples (n = 6) in each treatment and 17 treatments (Df = 16) in all. Significance codes: extremely significant 0 '***', highly significant p<0.001 '**', significant p<0.01 '*', slightly significant p<0.05 '..., and not significant p>0.05 'NS'.

Independent Variable	F-Value	n-Value	Significance Code
independent variable	i value	p vulue	Significance Code
Chlorophyll a+b	1.674	0.0676	
	1.000	0.0000	
Chlorophyll a	1.909	0.0303	*
Chlorophyll b	2.909	0.0007	***
Titratable Acidity (pH 7)	42.61	< 0.0001	***
Titratable Acidity (nH 7-	32.67	<0.0001	***
financie Actaity (pri)	52.01	-0.0001	
8.4)			
	16.60	0.0001	
Starch	16.68	< 0.0001	* * *
Glucose	7.189	< 0.0001	***
Fructose	6.279	< 0.0001	***
Sucrose	4 171	<0.0001	***
Sucrose	4.171	<0.0001	
NR Activity	26.22	< 0.0001	***
Nitrate	9.492	<0.0001	***
Ammonium	12.38	< 0.0001	***
Glyoxylic Acid	3.004	0.0001	***
NGD	24.6	-0.0001	ية باد يك
N:C Ratio	24.6	<0.0001	***
Percent C	1.39	0.1634	NS
Percent N	12.65	< 0.0001	***

Table 7: Ordinary one-way analysis (ANOVA) results of relative gene expression results with 6 samples (n = 6) in each treatment and 17 treatments (Df = 16) in all. Significance codes: extremely significant 0 '***', highly significant p<0.001 '**', significant p<0.01 '*', slightly significant p<0.05 '.'., and not significant p>0.05 'NS'.

Gene Names	F-Value	p-Value	Significance Code
ALMT	3.319	0.0001	***
PPC1	6.963	<0.0001	***
РРСК	3.460	< 0.0001	***
NR	10.290	< 0.0001	***
NiR	8.182	< 0.0001	***
GOGAT	4.176	< 0.0001	***
AS	3.885	< 0.0001	***
GDH460	3.773	< 0.0001	***
GDH201910	4.068	<0.0001	***
GS30900	1.950	0.0263	*
GS94700	13.490	< 0.0001	***





Figure 2: cladode length (cm) among treatments (n = 3). BT = measurements taken before applying treatments. Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 2: cladode width (cm) among treatments (n = 3). BT= measurements taken before applying treatments. Treatments are all modified Hoagland's solution with

varying amounts of nitrate and ammonium (mMol) and a deionized water

treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 3: cladode Thickness (mm) among treatments (mm) (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 4: New cladode number among treatments (n = 3) and before the treatment period (BT). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 5: Primary root number among treatments (n = 3). BT= measurements taken before applying treatments. Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 6: Average primary root Length (cm) among treatments (n = 3). BT= measurements taken before applying treatments. Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 7: Relative water content (%) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium

(mMol) and a deionized water treatment (diH_2O) control. Letters represent the result of Tukey's test (a = 0.05).



Chlorophyll Content

Figure 8: Chlorophyll content: Ca (blue), Cb (red), and Ca + Cb (green) content (mg/ml) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Error bars represent standard error.



Figure 9: Chlorophyll a + b content (mg/ml) among treatments n = 6. Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 10: Chlorophyll a content (mg/ml) among treatments n = 6. Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 11: Chlorophyll b content (mg/ml) among treatments n = 6. Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 12: Difference between dawn and dusk titratable tissue to pH 7 (malate equivalent) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized

water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 13: Difference between dawn dusk titratable tissue acidity from pH 7 to 8.4 (citrate equivalent) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 14: Starch content (mg/100ml of sample extract) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 15: Soluble sugars: glucose (blue), fructose (red), and sucrose (green) content (mg/100 ml sample extract) measured among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Error bars represent standard error. Negative values represent those calculated below zero by the standard linear equation for each respective sugar.



Figure 16: Glucose content (mg/mg of sample) among treatments (n = 6).

Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 17: Fructose content (mg/mg of sample) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 18: Sucrose content (mg/mg of sample) among treatments (n = 6).

Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 19: Nitrate reductase activity (nmoles NO₂ fixed/hour/gram fresh weight) among treatments (n = 3). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 20: Nitrate content (μ g/mg of sample) among treatments (n = 6).

Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 21: Ammonium content (μ Mol/mg of sample) among treatments (n = 3). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).


Figure 22: Glyoxylic acid content (μ Mol/mg of sample) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 23: N:C ratio among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 24: Percent carbon (%) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 25: Percent Nitrogen (%) among treatments (n = 6). Treatments are all modified Hoagland's solution with varying amounts of nitrate and ammonium (mMol) and a deionized water treatment (diH₂O) control. Letters represent the result of Tukey's test (a = 0.05).



Figure 26: Collective heatmap of N metabolism, and CAM related relative gene expression measured through RT-qPCR analysis among nitrate and ammonium treatments (mMol). Genes listed: aluminum activated malate transporter (ALMT), phosphoenolpyruvate carboxylase (PPC1), phosphoenolpyruvate carboxylase kinase (PPCK), nitrate reductase (NR), nitrite reductase (NiR), glutamate synthase (GOGAT), asparagine synthase (AS), glutamate dehydrogenase (GDH), and glutamine synthetase (GS). Relative expression of all genes is normalized to the 0+0 nitrate and ammonium treatment. The color scale represents actin and ubiquitin FPKM normalized log₂ transformed counts where blue indicates low expression, and red indicates high expression.



Figure 27: Relative expression of aluminum-activated malate transporter (ALMT) among nitrate and ammonium treatments (mM). Relative expression of ALMT in all treatments is normalized to average ALMT expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a=0.05). (n = 6)



Figure 28: Relative expression of phosphoenolpyruvate carboxylase (PPC1) among nitrate and ammonium treatments (mM). Relative expression of PPC1 in all treatments is normalized to the average PPC1 expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a = 0.05). (n = 6)



Figure 29: Relative expression of phosphoenolpyruvate carboxylase kinase (PPCK) among nitrate and ammonium treatments. Relative expression of PPCK in all treatments is normalized to the average PPCK expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a = 0.05). (n=6)



Figure 30: Relative expression of nitrate reductase (NR) among nitrate and ammonium treatments. Relative expression of NR in all treatments is normalized

to the average NR expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log_2 transformed counts. Letters represent the result of Tukey's test (a = 0.05). (*n* = 6)



Figure 31: Relative expression of nitrite reductase (NiR) among nitrate and ammonium treatments. Relative expression of NiR in all treatments is normalized to the average NiR expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a = 0.05). (n = 3)



Figure 32: Relative expression of glutamate synthase (GOGAT) among nitrate and ammonium treatments. Relative expression of GOGAT in all treatments is normalized to the average GOGAT expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a = 0.05). (*n*

= 3)



Figure 33: Relative expression of asparagine synthase (AS) among nitrate and ammonium treatments. Relative expression of AS in all treatments is normalized

to the average AS expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log_2 transformed counts. Letters represent the result of Tukey's test (a = 0.05). (*n* = 6)



Figure 34: Relative expression of glutamate dehydrogenase (GDH460) among nitrate and ammonium treatments. Relative expression of GDH460 in all treatments is normalized to the average GDH460 expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a = 0.05). (n = 6)



Figure 35: Relative expression of GDH (GDH201910) among nitrate and ammonium treatments. Relative expression of GDH201910 in all treatments is normalized to the average GDH201910 expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a = 0.05). (n = 6)



Figure 36: Relative expression of glutamine synthase (GS30900) among nitrate and ammonium treatments. Relative expression of GS30900 in all treatments is

normalized to the average GS30900 expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized \log_2 transformed counts. Letters represent the result of Tukey's test (a = 0.05). (*n*



Figure 37: Relative expression of glutamine synthetase (GS94700) among nitrate and ammonium treatments. Relative expression of GS94700 in all treatments is normalized to the average GS94700 expression in the 0+0 nitrate and ammonium treatment. All values represent the average actin and ubiquitin FPKM normalized log₂ transformed counts. Letters represent the result of Tukey's test (a = 0.05). (n = 6)

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Chapter 4

A life cycle inventory of estimated biogas and bioethanol production from an *Opuntia ficus-indica* field trial in the United States

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Abstract

Production of biofuels from *Opuntia spp.* could become a source of energy while having net negative CO₂ emissions. The global warming potential of agricultural production systems can be decreased with production of biogas or bioethanol from *Opuntia ficus-indica*. This study evaluates the production of biogas and bioethanol from *O. ficus-indica* in the United States using data from a previously published 5-year field trial in Logandale, Nevada. Biogas production of 13,004.29-26,877.85 Nm³ ha⁻¹ yr⁻¹ could be produced from plants receiving 716 mm year⁻¹, and 4082.46-8,532.19 Nm³ ha⁻¹ yr⁻¹ for plants receiving 300 mm ha⁻¹ year⁻¹ with a planting density of 1,418 plant ha⁻¹. Bioethanol production of 470.87 and 1783.32 Kg ha⁻¹ yr⁻¹ could be produced from plants at the same planting density receiving 300 and 716 mm yr⁻¹, respectively. The efficiency and bioenergy yields could easily be advanced if planting density were increased and fertilizer, transport, and harvesting methods were optimized.

Introduction

Opuntia ficus-indica (prickly pear cactus) is grown for the production of edible young cladodes, fruits (López, 1995), cattle feed (Mayer and Cushman, 2019), several secondary products, and most recently, biofuels (Yang et al., 2015). Central Mexico is the center of *Opuntia* spp. diversity (Omar et al., 2021), but it is currently grown in arid, semi-arid, and subtropical locations around the globe. Stem succulence, rectifier like roots, thick epidermal waxes, and utilization of crassulacean acid metabolism photosynthesis, a photosynthetic pathway in which carbon is assimilated at night, all confer an extremely high level of water use-efficiency (Nobel, 2003), while still maintaining competitive whole biomass production values between 2.4 to 47.3 Mg ha⁻¹ yr⁻¹ (Monjauze and Le Houérou, 1965; Garcia de Cortazar and Nobel, 1992; Dubeux et al., 2006; Guevara et al., 2011; Ramírez-Arpide et al., 2018; Neupane et al., 2021). Overall, *O. ficus-indica* has a recognized global economic value of \$2,520 million year⁻¹ globally (Davis et al., 2019).

As a cactus species, *Opuntia ficus-indica* operations can be established in the southwestern United States while requiring minimal water input (Neupane et al., 2021). *O. ficus*-indica can also cause increased CO₂ emissions from detrimental land use changes (Bautista-Cruz et al., 2018). In addition, *O. ficusindica* could be integrated with photovoltaic systems, where water run-off from cleaning of panels is enough to grow plants (Cushman et al., 2015). A 2.6% (w/v) cellulosic bioethanol conversion of cladode hydrosolates has been achieved (Kuloyo et al., 2014), but a minimal conversion efficiency of 4% (w/v) is considered standard for the success of a crop as a bioethanol source. As a biogas

source, O. ficus-indica has been demonstrated to contain 78% volatile solids (Calabrò et al., 2018), and a maximum methane production value of 604 NmL/gVSadded (Jigar et al., 2011). The addition of cow manure (Calabrò et al., 2018), and cow rumen fluid (Myovela et al., 2019) have both been shown to increase biogas production from O. ficus-indica. In addition, innoculation of microbes found degrading decaying cladodes in the soil of O. ficu-indica plantations have been shown to increase cladode digestion, due in part to the excretion of pectinase activity (Blair et al., 2021). Ethanol production from O. *ficus-indica* cladodes or fruits of 2.5 to 34.9 g L⁻¹ have been reported (Retamal et al., 1987; Kuloyo et al., 2014; de Souza Filho et al., 2016; do Nascimento Santos et al., 2016; Alencar et al., 2018; Pérez-Cadena et al., 2018), and biogas (methane) yields of 233 to 325 L kg⁻¹ have been reported from the anaerobic digestion of O. ficus-indica cladodes (Ramos-Suárez et al., 2014; Mason et al., 2015; Calabrò et al., 2018; Ramírez-Arpide et al., 2018; Valenti et al., 2018; Lueangwattanapong et al., 2020).

A complete life cycle assessment (LCA) and life cycle impact assessment (LCIA) of biogas production from *O. ficus-indica* under conventional farming methods using synthetic fertilizer, and an organic farming method using cow manure was conducted in Queretaro, Mexico (Ramírez-Arpide et al., 2018). The authors found that the organic farming scenario with nopal cladode transport, preprocessing by grinding, and mixing in 3:1 mix with cow manure, followed by anaerobic digestion in a covered lagoon anaerobic digester, and closed storage of the digestate for later use in inoculating the digestor had the highest energy return

on investment (amount of energy expended to produce a certain amount of net energy) value of 12.41, which was a 12% increase over the conventional farming base line scenario. In addition, global warming potential (a unit of emission of combined climate warming gases) was reduced by 2.3%. However, soil acidification and eutrophication potentials were increased under the organic farming scenarios. In total, in the organic farming plot, 417,130 Kg ha⁻¹ yr⁻¹ of biomass was produced with a planting density of 22,988 plant ha⁻¹, resulting in the production of 30,068.88 Mg yr⁻¹ of biogas.

In 2021, an *Opuntia spp.* field trial was conducted in the Logandale, Nevada to determine optimal irrigation inputs for production (Neupane et al., 2021). Within the irrigation treatments of 300, 407, and 716 mm year⁻¹ water input resulted in a combined average yield of 8.25, 11.16, and 15.52 Mg dry mass ha⁻¹ year⁻¹, respectively, for three different *Opuntia* spp. (Figure 1). Interestingly, *Opuntia cochinellifera,* which has longer oblong shaped cladodes, had lower occurrence of limb breakage due to wind, and accumulated the highest biomass after five years of growth. However, *Opuntia ficus-indica* was more responsive to water inputs and produced higher quantities of fruit with commercial grade fruit quality.

The objective of this study was to generate a life cycle inventory, as well as estimates of biogas and bioethanol production, using the inputs and data collected in the Logandale field trial. The resulting inventory was then used to make comparisons to the inventory and yields published by Ramírez-Arpide et al., 2018. These results were then used to performed complete life cycle assessment (LCA) of anerobic biogas and bioethanol fermentation of *O. ficus-indica* in the United States following the International Organization for Standardization framework (ISO) for direct comparisons against other bioenergy, and alternative fuel sources.

Methods

In 2021 our lab published the results of a five-year field trial of *Opuntia ficus-indica* in the United States (Neupane et al., 2021). This trial tested the above ground biomass production, fruit yield, and limb breakage in three different *Opuntia* spp. in combined precipitation and irrigation levels of 300 to 716 mm year⁻¹. A simple comparative life cycle inventory was constructed using the values from this study (Table 1) following the methods of (Ramírez-Arpide et al., 2018), which provided a full LCA of *O. ficus-indica* biogas production from an operation in Queretaro, Mexico.

The annual productivity of *O. ficus-indica* in the 300 to 716 mm year⁻¹ was calculated by taking the average biomass gained by *O. ficus-indica* between the 4th and 5th year (Figure 1) (Neupane et al., 2021) and multiplying it by the planting density, as we considered a coppicing operation in which only the cladodes produced after the 4th year of growth were used for biogas and ethanol production. A typical household-sized University of Nevada Cooperative Extension (UNCE) building was next to the field site, and grid electricity use of a typical American houshold was used (eia.gov). The Overton power district generates 25% of its monthly electricity from hyrdoelectic power from Hoover

damn (Figure 2). The remaining percent of energy generated from natural gas, coal, solar, and wind energy vary from month to month, but typically more than 50% of the electiricity generated is from natural gas. Fuel use was estimated by considering transport of harvested cladodes from the field to a transport truck approximated to be 0.2 km ha⁻¹ of ten trips up and down rows with a compact tractor (John Deere 1023E, 340 Kg carrying capacity, 14.48 Km/L diesel). The transport truck (GMC C7500 Box Truck, 4535.92 Kg carrying capacity, 4.67 Km/L diesel) was then used to transport harvested material to a theoretical covered lagoon anaerobic digester 50 km away for biogas production (Ramírez-Arpide et al., 2018), or a large-scale bioethanol fermentation plant (Kuloyo et al., 2014). The water inputs were the actual irrigation additions without including precipitation with the addition of supplemental water for inoculation of cladodes under digestion.

Theoretical biogas yield was first calculated by using published percent of volatile carbons in *O. ficus-indica* of 78% (Jigar et al., 2011) of 420 NmL/gVSadded from adding the raw substrate alone and a high methane yield 604 NmL/gVSadded with an acid pretreatment were used to estimate biogas generated in both low and high irrigation treatments per ha (Calabrò et al., 2018). Remaining effluent was estimated by calculating the percent effluent remaining from the methane generation in Ramírez-Arpide et al., 2018, and applying that percent to our own yields. Lastly, the estimated ethanol yield was calculated from a reported 2.6 % (w/v) (Kuloyo et al., 2014) obtained by fermentation of the *O. ficus-indica* cladode hydrolysate by hydrolysis and fermentation and simultaneous saccharification and fermentation procedures using *Kluyveromyces marxianus* and *Saccharomyces cerevisiae* at 35 and 40 °C.

Results

The Logandale field site produced an average biomass gain of approximately 12.77 and 40.23Kg/plant in the 300 to 716 mm year⁻¹ respectively, while Ramírez-Arpide et al., 2018 produced about 11.47 Kg/plant (Table 1). The Logandale field trial used about 3.5-fold more fertilizer than the conventional farming system in Ramírez-Arpide et al., 2018. Ramírez-Arpide et al., 2018 used slightly less grid electricity (28,271.1 MJ a⁻¹), and Mexico currently generates its electricity from natural gas, geothermal, hydro, nuclear, hard coal, fuel oil, and diesel sources (Figure 3). Ramírez-Arpide et al., 2018 used more water (2.07 Kg yr⁻¹) than even our highest irrigation treatment (0.580 Kg ha⁻¹ yr⁻¹), and similar diesel fuel use compared to our estimated value for the 716 mm yr⁻¹ treatment (222.2 vs. 230 L ha⁻¹ yr⁻¹).

The theoretical yield of biogas from *O. ficus-indica* biomass produced in Neupane et al., 2021 was 13,004.293-26,877.85 Nm³ ha⁻¹ yr⁻¹ for plants receiving 716 mm year⁻¹, and 4082.46-8,532.19 Nm³ ha⁻¹ yr⁻¹ for plants receiving 300 mm ha⁻¹ year⁻¹ (Table 1). This is more methane production per plant than reported in Ramírez-Arpide et al., 2018 (2.87-18.95 Nm³ yr⁻¹ plant⁻¹ vs. 1.308 Nm³ yr⁻¹ plant⁻¹), but less methane per hectare than the published 30,068.88 Nm³ ha⁻¹ yr⁻¹ value. Estimated effluent values were 2712.39 and 861.02 Kg ha⁻¹ yr⁻¹ for the 300 and 716 mm yr⁻¹, respectively. Lastly, estimated ethanol production for the Neupane et al., 2021 field trial was 470.87 and 1783.32 Kg ha⁻¹ yr⁻¹ for the 300 and 716 mm yr⁻¹, respectively.

Discussion

Several key differences were apparent between the Logandale field trial and that performed by Ramírez-Arpide et al., 2018 using a conventional farming system scenario. Mainly, the Logandale trial defined differences in production vs. irrigation instead of biogas production. Therefore, about 3.5-fold more nitrogen was applied to the plants to avoid limitations in growth due to nutrient availability. A much lower planting density was also used (22,988 vs. 1, 418 plants ha⁻¹) to insure that no intraspecific competition among individuals occurred and that no interference among irrigation blocks occurred at the field site. Initial field preparation required two rounds of tilling, a perimeter fence for herbivory prevention, and irrigation lines used for delivery of supplemental irrigation water and monthly all Purpose LiquaFeed® fertilizer N:P:K 12:4:8 (Scott's MiracleGro, Inc.). Weeds were controlled by hand weeding instead of herbicides, and no pesticides were used. In comparison, Ramírez-Arpide et al., 2018 site preparation involved initial harrowing and plowing and plants were grown for one year before reaching a productive stage for biomass estimates. Furthermore, dry inorganic fertilizer containing urea, triple superphosphate, and potassium chloride were used and weeds and pests were controled using atrazin and insecticides (unspecified), respectivly.

The measured biomass production per plant in the Logandale field trial was higher than the actual field values reported in Ramírez-Arpide et al., 2018. However, these values would likely be similar if planting density and fertilizer inputs were the same (Guimarães et al., 2020). Percentage of volatile carbons and methane production increased when a 3:1 mixture of *O. ficus-indica* with cow manure was used (Jigar et al., 2011). This finding is convenient when considering that *O. ficus-indica* makes for an excellent source of cattle fodder when combined with other feedstocks (Mayer and Cushman, 2019). Therefore, *O. ficus-indica* production and cattle ranching can be combined successfully into a single production system.

Field emissions from the soil were not measured in Neupane et al., 2021. The use of synthetic fertilizers in *O. ficus-indica* were shown to produce net negative methane emissions (-1,314.6 gCH₄ ha⁻¹ a⁻¹), but more N₂O emissions (636 gN₂O ha⁻¹ yr⁻¹) in the Ramírez-Arpide et al., 2018 trial compared to the organic scenario. The organic scenario in the same field used cow manure for fertilizer, producing a substantial amount of released methane (661.5g CH₄ ha⁻¹ yr⁻¹), but less N₂O (375.85 gCH₄ ha⁻¹ yr⁻¹). Using fertigation to directly apply fertilizer in irrigation lines has also been shown to reduce N₂O emissions compared to granular urea application in *Triticum aestivum* (wheat) and *Brassica napus* (canolla) (Chai et al., 2020), *Lycopersicon esculentum* (tomatoes) (Kennedy et al., 2013), and *Zea mays* (maize) (Tian et al., 2017). As such, the use of fertigation likely did not produce methane emissions, while also reducing N₂O emissions than if granular urea been applied instead.

The Logandale field trial benefitted from having information available on grid electricity sources on a local level from the Overton Power District (opd5.com) (Figure 2), whereas Ramírez-Arpide et al., 2018 used grid electricity sources from the whole country (Figure 3) (Itten et al., 2012). The majority of grid electricity from both operations was sourced from natural gas, which generates about 66 gCO₂/MJ in the United States (Venkatesh et al., 2011). Hydroelectricity makes up 25% of the grid electricity each month due to close proximity to Hoover dam. Hydroelectric power generates an estimated 23.6 gCO₂/MJ and 3.0 gCH₄/MJ as the flow of water is slowed, and organic matter builds up and degrades anerobically (Hertwich, 2013). Solar and wind electricity generation vary from month to month, and must be supplemented by natural during periods when renewable energy production is limited. Southern Nevada recieves a high amount of solar radiation, and a minimum of 9.7 gCO₂/MJ can be achieved with solar panels (Liu et al., 2015). Advancement of battery technology has also helped to increase the efficiency of wind gereration, and 5 gCO₂/kWh can be achieved with wind power (Alsaleh and Sattler, 2019). Electricity generated in Overton power district most certainly coresponded with less CO₂ emissions than the Mexican power grid (Figure 3) as it uses nuclear (3.3) gCO₂/MJ), hard coal (303.8 gCO₂/MJ), fuel oil (267 gCO₂/MJ) (Santoyo-Castelazo et al., 2011), and diesel in addition to natural gas, wind, and hydroelectricity.

Estimated diesel used for tranportation was higher at the Logandale field site, and the efficiency could have been improved by using a tractor with a larger carrying capacity although the compact tractor was convinient for navigating narrow rows for research purposes. The CO_2 emmissions from the transport truck could be mitigated by primarily using rail ways for transport of cladodes to a centralized digestor, as rail transport has the lowest associated CO_2 emmissions (Horvath, 2006), although this would require strategic planning of location along a rail line. Current battery powered trucks are a promising form of transportation in regards to CO_2 emmissions, but do not significantly improve life-cycle emissions or costs compared to diesel without generating all the energy for operation and manufacturing with biofuels (Sen et al., 2017).

Both the life cycle inventory created here and in Ramírez-Arpide et al., 2018 did not consider the impact of human labor as an input, and including human labor has been shown to have about a 10.2% increase in embodied energy cost esimates in agricultural production pathways (Rocco and Colombo, 2016). In addition, human labor costs are much cheaper in Mexico (Roka and Guan, 2018) and would likely be a major barrier in the competitive biofuel production of *O*. *ficus-indica* in the United States. As a solution, mechanization of *O*. *ficus-indica* harvesting for biofuel generation is concievable with typical farm equipment where the entire above ground biomass is harvested, but specialized machinery would be needed for the coppicing operations outlined here.

In conclusion, these life cycle inventory estimates indicate that theoretical biogas production from *O. ficus-indica* in Logandale, Nevada is comparable to that of actual biogas production values in Mexico, but without the use of herbicides or pesticides. *O. ficus-indica* agriculture, in combination with other

renewable energy sources, could help the United States meet demands for energy sources with reduced net CO₂ emissions. The next step of this study is to follow the methods of Ramírez-Arpide et al., 2018, in creating a complete LCA of *O*. *ficus-indica* following ISO guidelines (ISO), using OpenLCA software (GreenDelta, 2017), or a similar software (Silva et al., 2017), and the ecoinvent database (Wernet et al., 2016).

Tables

Table 1: A life cycle inventory (LCI) for *Opuntia ficus-indica* grown in an

irrigation treatment in Logandale, NV, United States (Neupane et al., 2021).

Parameter	Units	Quantity
Productivity		
Planting Density	Plants ha ⁻¹	1,418
High Irrigation <i>Ofi</i> productivity (716 mm yr ⁻ ¹)	Kg d.w. ha ⁻¹ yr ⁻¹	57,050
Low Irrigation <i>Ofi</i> productivity (300 mm yr ⁻ ¹)	Kg d.w. ha ⁻¹ yr ⁻¹	18,110
High Irrigation <i>Ofi</i> productivity (716 mm yr ⁻ ¹)	Kg f.w. ha ⁻¹ yr ⁻¹	298848
Low Irrigation <i>Ofi</i> productivity (300 mm yr ⁻¹)	Kg f.w. ha ⁻¹ yr ⁻¹	221534
Inputs		
Urea	Kg ha ⁻¹ yr ⁻¹	875.75
P ₂ O ₅	Kg ha ⁻¹ yr ⁻¹	291.91
K ₂ O	Kg ha ⁻¹ yr ⁻¹	583.83
Mn	Kg ha ⁻¹ yr ⁻¹	3.65
Chelated Mn	Kg ha ⁻¹ yr ⁻¹	3.65
Zn	Kg ha ⁻¹ yr ⁻¹	3.65
Chelated Zn	Kg ha ⁻¹ yr ⁻¹	3.65
Grid Electricity	MJ ha ⁻¹ yr ⁻¹	38,574 (Average household eia.gov)
Diesel (716 mm yr ⁻¹)	L ha ⁻¹ yr ⁻¹	230
Diesel (300 mm yr ⁻¹)	L ha ⁻¹ yr ⁻¹	170
Water (716 mm yr ⁻¹)	mm ha ⁻¹ yr ⁻¹	580
Water (300 mm yr ⁻¹)	mm ha ⁻¹ yr ⁻¹	164

Theoretical Yield		
Biogas (716 mm yr ⁻¹)	Nm ³ ha ⁻¹ yr ⁻¹	1,300-26,877 (Calabrò et al., 2018; Jigar et al., 2011)
Biogas (300 mm yr ⁻¹)	Nm ³ ha ⁻¹ yr ⁻¹	4082-8,532 (Calabrò et al., 2018; Jigar et al., 2011)
Effluent (716 mm yr ⁻¹)	Kg ha ⁻¹ yr ⁻¹	2712 (Ramírez-Arpide et al., 2018)
Effluent (300 mm yr ⁻¹)	Kg ha ⁻¹ yr ⁻¹	861 (Ramírez-Arpide et al., 2018)
Ethanol (716 mm yr ⁻¹)	Kg ha ⁻¹ yr ⁻¹	1483 (Kuloyo et al., 2014)
Ethanol (300 mm yr ⁻¹)	Kg ha ⁻¹ yr ⁻¹	470.87 (Kuloyo et al., 2014)
Figures



Figure 1: *Opuntia spp.* above ground fresh weight (FW) productivity under varying irrigation inputs over 5 years in Logandale, Nevada. Bars represent the average above ground FW biomass (Kg) of cladodes *Opuntia ficus-indica* (OFI: light blue, blue, dark blue), *Opuntia streptacantha* (OS: light red, red, dark red), and *Opuntia cochenillifera* (OC: light green, green, dark green). Error bars represent standard error (n = 4). Data sourced from Neupane et al. 2021.



Figure 2: Nevada Overton power district energy source break down. (source: eia.gov and opd5.com)



Figure 3: Contribution of different fuels to the electricity mix in Mexico. Figure adapted from Ramírez-Arpide et al., 2018

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Chapter 5

Concluding Remarks

Of the three photosynthetic pathways, CAM confers the highest water-use efficiency (Borland et al., 2009), and often occurs with several co-adapted traits that enhance tolerances to other types of abiotic stress (Niechayev et al., 2019). As a desert-adapted species, Opuntia ficus-indica is an obligate CAM species with a remarkable growth rate, tasty fruits, vegetable like young cladodes, cosmetic seed oil, and provides feed for livestock, and shows comparatively high biogas production values (Lueangwattanapong et al., 2020). In a drying world with ever decreasing ground water stores (Naumann et al., 2018), O. ficus-indica and other economically important CAM species (Davis et al., 2019) may prove to be vital in maintaining food, energy, and economic security in regions that regularly experience drought. In chapter 2 young tissues in O. ficus-indica were shown to use CAM early in development; in chapter 3 plant growth, biochemistry, and N and CAM gene expression is affected by variation in N source; and in chapter 4 potential biofuel production from O. ficus-indica in the United States was estimated.

In chapter 2, *O. ficus-indica* seedlings were shown to use CAM photosynthesis early in development, even under well-watered conditions. Seedlings also continued to use CAM photosynthesis to fix CO_2 even when the soil was completely dried out. These findings demonstrated that C_3 photosynthesis does not play a major role in seedling development. However, some daytime fixation of CO_2 does occur at dawn and dusk, especially when light intensity was increased. In contrast, *O. eliator* (Winter and Holtum, 2011), a tropical epiphyte species was shown to perform C₃ photosynthesis early on with none of the defined CAM phases, and fix CO₂ predominantly during the light period unless under drought conditions. Collectively, these findings demonstrate that the diversity in photosynthetic plasticity in the *Opuntia* genus with tropical species exhibiting more facultative CAM, and xeric species showing more obligate CAM.

O. ficus-indica daughter cladodes had previously been demonstrated to act as sink tissues before beginning in CAM photosynthesis (Wang et al., 1998). Our results corroborated these findings, as the isotopic ratios were indicative of typical CAM fixation no matter what size the cladodes were, nocturnal tissue acidity began to build-up early on, and CO_2 assimilation occurred primarily during the dark period. However, this was in contrast with the results of (Winter et al., 2008) who measured higher net CO_2 assimilation during the light period. The same results were observed when *O. ficus-indica* cladodes were measured under optimal conditions in a growth chamber instead of in a greenhouse. Higher daytime net CO_2 assimilation observed in the growth chamber is likely due to constant light conditions that cause a higher fixation of CO_2 before the lights turn off, and after they turn on *vs.* the greenhouse in which the sun gradually sets and rises, limiting the amount of CO_2 assimilation during twilight. The greenhouse conditions may better represent how *O. ficus-indica* assimilates CO_2 in the field.

In chapter 3, under varying nitrate and ammonia concentrations in sand culture, statistically significant differences in *O. ficus-indica* growth, chlorophyll

content, tissue acidity, soluble sugars, nitrate reductase activity, nitrate and ammonia content, glyoxylic acid content, N:C ratio and relative expression of genes involved N metabolism, and CAM activity were measured. Our deionized water treatment demonstrated that when nutrients are limited growth is hindered, tissue acidity, and starch content is increased, and soluble sugars are no longer present. Within deionized water treatment, cladodes also showed increased steady-state mRNA expression of the CAM-related genes encoding aluminumactivated malate transporter (ALMT) and phosphoenolpyruvate carboxylase (PEPC) and higher steady-state mRNA expression of the nitrogen metabolismrelated genes including glutamine oxoglutarate aminotransferase (Fd-GOGAT), glutamate dehydrogenase (NADH-GDH460), glutamine synthetase 2 (GS2), but lower expression of asparagine synthetase (AS) than nutrient supplied cladodes. Altogether, these results suggest that when nutrients are limited, *Opuntia ficusindica* converts soluble sugars into starch, builds up stored organic acids, and enhances expression of CAM-related genes.

In general, the responses in growth, biochemistry, and gene expression reflected that growth and metabolism were enhanced when more nitrogen was present regardless of whether *O. ficus-indica* was given nitrate or ammonium. (Vázquez et al., 2000) et al. supported this evidence as *O. ficus-indica* cladodes were shown to take up N regardless of being given only nitrate or ammonia, and only had slight increases in N uptake when given ammonium alone. *O. ficusindica* field operations can be successful with either nitrate (Nerd and Nobel, 1995), ammonium (Nkoi et al., 2021), or urea (Galizzi et al., 2004), as long as soil pH is kept balanced with N application. In retrospect, it would have been useful to collect samples at multiple timepoints with variation in nitrate and ammonium, although our sample size was already cumbersome for a single timepoint (~112 samples * 34 independent variables measured). Therefore, repeating the experiment with fewer nitrate and ammonia treatments and/or singling out desired measurements from our methodology, and adding more time points might reveal trends overtime in relation to diversity of N source.

In Chapter 4, beginnings of a life cycle assessment for biogas and bioethanol production from *O. ficus-indica* field trial in the United States (Neupane et al., 2021) was presented in the form of a life cycle inventory. Our inventory mirrored that of a complete life cycle assessment from an *O. ficusindica* trial in Mexico (Ramírez-Arpide et al., 2018) in order to have the ability to make comparisons between the two trials. Clearly, the United States irrigation trial was not designed with biogas generation in mind, and therefore, had a much higher application of fertilizer, and space between plants. Having plants in optimal nutrient and spacing conditions resulted in higher yields on a per plant basis, but the higher planting density in the Mexican trial resulted in higher overall yield.

Several advantages became apparent when comparing an operation in the United States *vs.* Mexico. First, being that grid electricity in Southern Nevada is typically produced with more renewable energy sources and less carbon emissions than that of Mexico. Second, the Logandale field trial, herbicides and insecticides were not needed. Both of these advantages would decrease the net carbon release associated with biogas production from O. ficus-indica in the United States.

Appearance of *Cactoblastis* (Zimmermann et al., 2004) or *Cochineal* (Moran and Cabby, 1979; Vigueras et al., 2007) pests could conceivably happen at any time and require insecticides to be used. *Cochineal* is a problem in our ongoing field trials in Parlier, CA. As a disadvantage, field labor in the United States is much more expensive than Mexico, and competitive unit biogas production per dollar from *O. ficus-indica* would likely require mechanized harvesting.

Conclusion

Opuntia ficus-indica is a highly productive CAM species that can improve food, feed, and fuel security in drought prone regions. As a drought adapted species it predominantly uses CAM throughout development. In regard to N source preferences, *O. ficus-indica* grows slightly better when given slightly more nitrate than ammonium. Finally, estimated biofuel production from *O. ficusindica* in the United States is comparable to that of actual yields in Mexico, and biogas production from *O. ficus-indica* could be used to produce biofuels with minimal CO₂ emissions associated with land use changes or effects on current food supply.

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Supplemental figures for chapter 2 Seedling 1 - PAR 100 - Watered



Figure 1: 24-hour gas exchange measurements of *O. ficus-indica* seedling 1 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20°C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 2: 24-hour gas exchange measurements of *O. ficus-indica* seedling 2 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 3: 24-hour gas exchange measurements of *O. ficus-indica* seedling 3 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 4: 24-hour gas exchange measurements of *O. ficus-indica* seedling 4 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes. This seedling had chlorosis from tightly packed soil and was not used to calculate the AUC for net gas exchange of well-watered plants in PAR 100.



Figure 3: 24-hour gas exchange measurements of *O. ficus-indica* seedling 5 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.

Figure 6: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 1 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.

Figure 7: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 2 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 8: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 3 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 9: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 4 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions. This seedling had chlorosis from tightly packed soil and was not used to calculate the AUC for net gas exchange of well-watered plants in PAR 100.



Figure 40: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 5 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 11: The combined diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in all seedlings grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 12: 24-hour gas exchange measurements of *O. ficus-indica* seedling 1 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes. Each measurement was taken after allowing the soil to completely dry out for one day.

Seedling 2 - PAR 100 - Drought



Figure 13: 24-hour gas exchange measurements of *O. ficus-indica* seedling 2 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes. Each measurement was taken after allowing the soil to completely dry out for one day.



Figure 14: 24-hour gas exchange measurements of *O. ficus-indica* seedling 4 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows. The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes. Each measurement was taken after allowing the soil to completely dry out for one day. This seedling had chlorosis from tightly packed soil and was not used to calculate the AUC for net gas exchange of drought plants in PAR 100.

Seedling 5 - PAR 100 - Drought **6** Pad Length (cm) 0.0 An (μ mol CO₂ m⁻² s⁻¹) 3.5 7.0 2 0 6:00:00 OF 0:00:00 12:00:00 00:00 0n 24:00:00

Time (h:m:s)

Figure 15: 24-hour gas exchange measurements of *O. ficus-indica* seedling 5 grown at a PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes. Each measurement was taken after allowing the soil to completely dry out for one day.



Figure 16: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 1 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in drought. This seedling had chlorosis from tightly packed soil and was not used to calculate the AUC for net gas exchange of well-watered plants in PAR 100.



Figure 17: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 2 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in drought.


Figure 18: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 4 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in drought. This seedling had chlorosis from tightly packed soil and was not used to calculate the AUC for net gas exchange of well-watered plants in PAR 100.



Figure 19: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in seedling 5 grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in Drought.



Figure 20: The combined diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in all seedlings grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in drought.



Figure 21: Measured length (cm) of *O. ficus- indica* seedling 1 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the gas exchange sample period (days after germination). Grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 22: Measured length (cm) of *O. ficus- indica* seedling 2 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the

gas exchange sample period (days after germination). Grown at PAR of 100 μmol $m^{-2}~s^{-1}$ at 20 °C in well-watered conditions.



Figure 23: Measured length (cm) of *O. ficus- indica* seedling 3 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the gas exchange sample period (days after germination). Grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 24: Measured length (cm) of *O. ficus- indica* seedling 4 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the gas exchange sample period (days after germination). Grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 25: Measured length (cm) of *O. ficus- indica* seedling 1 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the

gas exchange sample period (days after germination). Grown at PAR of 100 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 5: 24-hour gas exchange measurements of *O. ficus-indica* seedling 6 grown at a PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows. The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 27: 1st set of 6 24-hour gas exchange measurements of *O. ficus-indica* seedling 6 grown at a PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 28: 2^{nd} set of 6 24-hour gas exchange measurements of *O. ficus-indica* seedling 6 grown at a PAR of 300 µmol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (µmol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 29: 3^{rd} set of 6 24-hour gas exchange measurements of *O. ficus-indica* seedling 6 grown at a PAR of 300 µmol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (µmol CO₂ m⁻² S⁻¹) was logged every 5 minutes.

Seedling 6 - PAR 300 - 4rth Set



Figure 30: 4rth set of 6 24-hour gas exchange measurements of *O. ficus-indica* seedling 6 grown at a PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 31: 24-hour gas exchange measurements of *O. ficus-indica* seedling 7 grown at a PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.

Seedling 7 - PAR 300



Figure 32: 24-hour gas exchange measurements of *O. ficus-indica* seedling 8 grown at a PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 33: 24-hour gas exchange measurements of *O. ficus-indica* seedling 9 grown at a PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions as the central cladode grows (Dark blue to light blue to light red to dark red). The assimilation rate (μ mol CO₂ m⁻² S⁻¹) was logged every 5 minutes.



Figure 34: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in *O. ficus-indica* seedling 6 grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 35: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in *O. ficus-indica* seedling 7 grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 36: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in *O. ficus-indica* seedling 8 grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 37: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (μ mol m⁻²) *vs.* cladode length (cm) in *O. ficus-indica* seedling 9 grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 38: Measured length (cm) of *O. ficus-indica* seedling 6 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the gas exchange sample period (days after germination). Grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 39: Measured length (cm) of *O. ficus-indica* seedling 7 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the

gas exchange sample period (days after germination). Grown at PAR of 300 μmol $m^{-2}~s^{-1}$ at 20 °C in well-watered conditions.



Figure 40: Measured length (cm) of *O. ficus-indica* seedling 8 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the gas exchange sample period (days after germination). Grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 41: Measured length (cm) of *O. ficus-indica* seedling 9 central cladode (blue dots and line) and combined cotyledons (orange dots and line) within the gas exchange sample period (days after germination). Grown at PAR of 300 μ mol m⁻² s⁻¹ at 20 °C in well-watered conditions.



Figure 42: 24-hour gas exchange measurements of *O. ficus-indica* daughter cladode 1 in the custom-built chamber as the cladode grows (Dark blue to light blue to light red to dark red) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night. The Δ CO2 (ppm) is equal to the sample chamber subtracted by the reference chamber and was logged every 10 minutes.



Figure 43: 24-hour gas exchange measurements of *O. ficus-indica* daughter cladode 2 in the custom-built chamber as the cladode grows (Dark blue to light blue to light red to dark red) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night. The Δ CO2 (ppm) is equal to the sample chamber subtracted by the reference chamber and was logged every 10 minutes.



Figure 44: 24-hour gas exchange measurements of *O. ficus-indica* daughter cladode 3 in the custom-built chamber as the cladode grows (Dark blue to light blue to light red to dark red) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night. The Δ CO2 (ppm) is equal to the sample chamber subtracted by the reference chamber and was logged every 10 minutes.



Figure 45: 24-hour gas exchange measurements of *O. ficus-indica* daughter cladode 4 in the custom-built chamber as the cladode grows (Dark blue to light blue to light red to dark red) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night. The Δ CO2 (ppm) is equal to the sample chamber subtracted by the reference chamber and was logged every 10 minutes.



Time (h:m:s)

Figure 46: 24-hour gas exchange measurements of *O. ficus-indica* daughter cladode 5 in the custom-built chamber as the cladode grows (Dark blue to light blue to light red to dark red) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night. The Δ CO2 (ppm) is equal to the sample chamber subtracted by the reference chamber and was logged every 10 minutes.



Figure 47: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladode 1 grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 48: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladode 2 grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 49: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladode 3 grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 50: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladode 4 grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.

Daughter Pad 4



Figure 51: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladode 5 grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 52: Combined diel (blue circle), light (yellow square), and dark (black triangle) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladodes grown under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 53: Measured length (cm) of *O. ficus-indica* daughter cladode 1 (blue dots and line) within the gas exchange sample period (days in chamber) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 54: Measured length (cm) of *O. ficus-indica* daughter cladode 2 (blue dots and line) within the gas exchange sample period (days in chamber) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 55: Measured length (cm) of *O. ficus-indica* daughter cladode 3 (blue dots and line) within the gas exchange sample period (days in chamber) under standard

greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 56: Measured length (cm) of *O. ficus-indica* daughter cladode 4 (blue dots and line) within the gas exchange sample period (days in chamber) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.



Figure 57: Measured length (cm) of *O. ficus-indica* daughter cladode 5 (blue dots and line) within the gas exchange sample period (days in chamber) under standard greenhouse conditions with natural light at approx. 1,100-1,500 μ mol m⁻² s⁻¹ and temperature at 28-32 °C day/17-18 °C night.


Figure 58: 24-hour gas exchange measurements of *O. ficus-indica* daughter cladode in the custom-built chamber as the cladode grows (dark blue to light blue to light red to dark red) within a Conviron growth chamber with a measured PAR at the canopy level of 367 μ mol m⁻² s⁻¹ and temperature at 25 °C day and night. The Δ CO₂ (ppm) is equal to the sample chamber subtracted by the reference chamber and was logged every 10 minutes.



Figure 59: The diel (blue line), light (yellow line), and dark (black line) net assimilated CO₂ (ppm) *vs.* cladode length (cm) in *O. ficus-indica* daughter cladode on an individual growing within a Conviron growth chamber with a measured PAR at the canopy level of 367 μ mol m⁻² s⁻¹ and temperature at 25 °C day and night.



Figure 60: Measured length (cm) of *O. ficus-indica* daughter cladode (blue dots and line) within the gas exchange sample period (days in chamber) within a Conviron growth chamber with a measured PAR at the canopy level of 367 μ mol m⁻² s⁻¹ and temperature at 25 °C day and night.

Germination with and without sucrose



Figure 61: Timing and frequency of germination events in *Opuntia ficus-indica* plated in either 3.0% sucrose, 0.6% plant agar, 0.5% Murashige & Skoog medium with Gamborg's B5 vitamins, and a 10mM Gibberellic acid concentration or the same media containing no sucrose. 94 out of 150 seeds germinated in sucrose, and 95 out of 150 seeds germinated in no sucrose. There were no germination events after 11 days.





Figure 62: Observed *O. ficus-indica* Seedling phenotypes included those with roots emerging from where the cladodes made contact with the hypocotyl (A), tricotyledonous seedling (B), polyembryonic seedlings (C), and di-cladode individuals (D). None of these abnormal phenotypes were used in the experimentation in this study.

Curriculum vitae

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EDUCATION

University of Nevada, Reno | Reno, NV **Ph.D. in Biochemistry 2017-present**

Ohio University | Athens, OH M.S. in Environmental Studies 2014-2016

California State University of San Bernardino | San Bernardino, CA **B.S. in Biology 2010-2014**

GRANTS, AWARDS, AND STIPENDS

USDA NIFA Grant awarded 2018 Second place at Ohio University's student research expo 2016 Ohio University Graduate Assistantship Stipend 2014 CSUSB Student Research Competition Award 2014 Dean's List 2013 & 2014 OSR Stipend Awarded 2013 ASI Materials Grant Awarded 2012 ASI Travel Grant Awarded 2012

TEACHING EXPERIENCE

University of Nevada, Reno | Reno, NV **Biochemistry Teaching Assistant** 2017 Led discussion sections, and graded exams.

California State University of San Bernardino | San Bernardino, CA Biology Tutor 2011-2014 Tutored cellular biology courses and graded exams

RELATED EXPERIENCE

University of Nevada, Reno | Reno, NV

2017-present

Research Assistant in Dr. John Cushman's Biochemistry Lab

Conducting research related to the development, transformation, and productivity of the potential biofuel feedstock *O. ficus-indica*.

Ohio University | Athens, OH 2014-2016 Graduate Assistant in Dr. Sarah Davis' Bioenergy and Ecology Lab Assisted with the construction and operation of an experimental anaerobic digester, farming of potential biofuel crops, and lab and

field experimentation of the potential biofuel feedstock Agave americana.

California State University of San Bernardino | San Bernardino, CA 2011-2014

Research Assistant in Dr. John Skillman's Plant Eco Physiology Lab Provided assistance in the study of seasonal variation in microclimate, plant phenology, and leaf physiology for *Carpobrotus edulis* and *Dudleya edulis* in the San Clemente coastal bluff vegetation, requiring techniques used in Chemistry, Biology, and Statistics.

WORKS AND PUBLICATIONS

- Niechayev NA, Mayer JA, Cushman JC. (2021) Developmental dynamics of crassulacean acid metabolism (CAM) in *Opuntia ficus-indica*, an obligate CAM species. *in preparation*.
- Niechayev NA, Pereira PN, & Cushman JC. (2021) The nitrate vs.. ammonium preference of *Opuntia ficus-indica*: a sand culture snapshot. *in preparation*.
- Niechayev NA, & Cushman JC. (2021). The economic and environmental viability of *Opuntia ficus-indica* in the southwestern United States. *in preparation*.
- Ferris C, King N, Niechayev NA, Mayer JA, & Cushman JC. (2021) An optimized Biolistic[®] and Agrobacterium mediated transformation of Opuntia ficus-indica. in preparation.
- Angres G, Niechayev NA, & Cushman JC. (2021) A field validated cladode area index (CAI) model for *Opuntia spp*. indirect measurements. *in preparation*.
- Neupane D, Mayer JA, Niechayev NA, Bishop CD, & Cushman JC. (2021). Five-year field trial of the biomass productivity and water input response of cactus pear (Opuntia spp.) as a bioenergy feedstock for arid lands. *GCB Bioenergy*, 13(4), 719-741. DOI: 10.1111/gcbb.12805
- Pereira, PN, Niechayev, NA, Blair BB., & Cushman JC. (2021). Climate change responses and adaptations in Crassulacean Acid metabolism (CAM) plants. In *Photosynthesis, Respiration, and Climate Change* (pp. 283-329). Springer, Cham. DOI: 10.1007/978-3-030-64926-5 10
- Davis SC, Simpson J, Gil-Vega KDC, Niechayev NA, Tongerlo EV, Castano NH, Dever LV, & Búrquez, A. (2019). Undervalued potential of crassulacean acid metabolism for current and future agricultural production. *Journal of Experimental Botany*, 70(22), 6521-6537. DOI:10.1093/jxb/erz223
- Niechayev NA, Pereira PN, & Cushman JC. (2019). Understanding trait diversity associated with crassulacean acid metabolism (CAM). Current opinion in plant biology, 49, 74-85. DOI: 10.1016/j.pbi.2019.06.004

- Niechayev NA, Jones AM, Rosenthal DM, & Davis SC. (2019). A model of environmental limitations on production of Agave americana L. grown as a biofuel crop in semi-arid regions. *Journal of Experimental Botany*, 70(22), 6549-6559. DOI:10.1093/jxb/ery383
- Davis SC, Kuzmick ER, Niechayev NA, & Hunsaker DJ. (2017). Productivity and water use efficiency of Agave americana in the first field trial as bioenergy feedstock on arid lands. GCB Bioenergy. 9:314-325. DOI: 10.1111/gcbb.12324

POSTERS AND TALKS

- Niechayev NA, Mayer JA, Henitz C, Bishop C, Neupane D, Pereira P, Angres G, Faris C, King N, Enriquez, Petrusa L, Cushman JC (2021). Development of *Opuntia ficusindica* for commercial production in the United States. International Virtual Mini-Symposium. Cactacea: Phylogenetics, Evolution and Conservation in the Genomics Era. Phoenix, AZ. September 1-3, 2021.
- Neupane D, Niechayev N, Mayer JA, Bishop CD, Cushman JC. Biomass productivity of prickly pear cactus (*Opuntia* spp.) under different irrigation regimes. American Society of Plant Biologists annual meeting, Washington, D.C. July 27-31, 2020.
- Niechayev N, Mayer JA, Cushman JC. Developmental dynamics of crassulacean acid metabolism (CAM) in *Opuntia ficus-indica*. American Society of Plant Biologists annual meeting, San Jose, CA. August 4-7, 2019.
- Niechayev N, Mayer JA, Cushman JC. Developmental dynamics of crassulacean acid metabolism (CAM) in *Opuntia ficus-indica*. Biology of CAM Plants, Desert Botanical Garden, Phoenix, AZ. March 18-22, 2018.
- Niechayev N, Mayer JA, Yim,WC, Cushman JC. Developmental dynamics of crassulacean acid metabolism (CAM) in *Opuntia ficus-indica*. 26th Plant and Animal Genome Conference, San Diego, CA. January 12-17, 2018.
- Niechayev NA., Beechko AN., Skillman JB. Seasonal variation in microclimate, plant phenology, and leaf physiology for *Carpobrotus edulis* and *Dudleya edulis* in the San Clemente coastal bluff vegetation. Conference paper and poster session presented at the 34th New Phytologist Symposium on Systems Biology and Ecology of CAM Plants, Tahoe City, *CA*. July 14, 2014.

RELEVANT SKILLS

Completed an in-person factory certified 3-day LI-COR LI-6400XT portable photosynthesis training course, and highly experienced with the LI-COR 6800 portable photosynthesis machine.

Experienced with R, R-studio, PRISM and Matlab statistical programs.

Proficient with Microsoft Office.

Hands on research experience with equipment and techniques involved in field and lab experimentation such as:

- Soil microbe Gas Chromatography
- LI-COR LI-6400 portable photosynthesis systems and light sensors

- Tissue sample titrations
- o RT-QPCR and relative expression analysis of target genes
- Isotopic mass spectrometric analysis
- Water potential determination via pressure bomb
- o DNA, RNA, protein, and specific metabolite extraction from plant tissues
- o Colorimetric assays
- o LECO FP928 combustion analyzer operation for N:C ratio determination
- Field trial design and experimentation
- Indirect measurement modeling
- Agricultural/environmental production modeling
- Tissue culture and greenhouse techniques for restoring and maintaining collections
- \circ Seed germination
- o Flow cytometry and microscopy for ploidy counting
- Plasmid cloning and plant transformation (*Agrobacterium* and biolistic) protocols
- Oxygen electrodes for photosynthetic and dark respiration rate measurements
- Collection and analysis of soil moisture, texture, and cation exchange capacity
- Growth and life/size stage censuring
- Slide Prep and Microscopy

Possess a Nevada Driver's License and completed a California defensive driving course for the purpose of safely transporting hazardous chemicals.

Years of hands-on experience restoring, maintaining, and operating tractors, automobiles, motorcycles, and small engines.

Avid gardener.

World Tae Kwon Doe Federation certified 1st dan black belt.