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**Rhizodeposition and biotic interactions in the rhizosphere of  
*Phaseolus vulgaris* L. and *Hordeum vulgare* L.**

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Stuttgart, den 11.01.2008

Susan Haase

# Contents

1	Summary .....	8
2	Zusammenfassung .....	11
3	General Introduction .....	15
3.1	Release of organic compounds by living plant roots .....	15
3.2	Elevated atmospheric CO <sub>2</sub> concentration.....	16
3.2.1	Effects of elevated CO <sub>2</sub> on belowground processes - rhizodeposition .....	18
3.2.2	Effects of elevated CO <sub>2</sub> on soil microbes.....	19
3.3	Belowground herbivory by plant-parasitic nematodes.....	22
3.4	Collection of root exudates .....	23
4	Thesis Outline .....	25
5	Elevation of atmospheric CO <sub>2</sub> and N-nutritional status modify nodulation, nodule-carbon supply, and root exudation of <i>Phaseolus vulgaris</i> L. ....	27
5.1	Abstract .....	28
6	Local response of bacterial densities and enzyme activities to elevated atmospheric CO <sub>2</sub> and different N supply in the rhizosphere of <i>Phaseolus vulgaris</i> L. ....	29
6.1	Abstract .....	30
7	Responses to iron limitation in <i>Hordeum vulgare</i> L. as affected by the atmospheric CO <sub>2</sub> concentration .....	31
7.1	Abstract .....	32
8	Low-level herbivory by root-knot nematodes ( <i>Meloidogyne incognita</i> ) modifies root hair morphology and rhizodeposition in host plants ( <i>Hordeum vulgare</i> ) .....	33
8.1	Abstract .....	34
9	General Discussion and Concluding Remarks .....	35
9.1	Effects of elevated atmospheric CO <sub>2</sub> concentration.....	35
9.2	Effects of low-level herbivory by plant-parasitic nematodes.....	36
10	References .....	38

Curriculum vitae.....	57
Publications and Presentations .....	59
Acknowledgements .....	63

## List of Figures

- Fig. 3.1:** Localized collection of root exudates from soil grown plants by use of sorption media (filter papers)..... 24
- Fig. 3.2:** (a) PVC mini-rhizotron (rhizobox) consisting of a corpus with irrigation holes and a perspex front lid. (b) Rhizobox holder for fixing rhizoboxes during the culture period..... 24

## 1 Summary

Biochemical processes at the soil-plant interface are largely regulated by organic and inorganic compounds released by roots and microorganisms. Several abiotic and biotic factors are suspected to stimulate rhizodeposition and, thus, contribute to enriching of the rhizosphere with plant-derived compounds. This thesis focused on the effects of two factors, (i) the elevation of atmospheric CO<sub>2</sub> concentration accompanied by nutrient limitation in the soil and (ii) low-level root infestation by plant-parasitic nematodes, on the quantity and quality of rhizodeposits with consequences for plant-nutrient acquisition and plant-microbial interactions in the rhizosphere. Experiments were largely conducted in mini-rhizotrons, which allowed a localized collection of rhizodeposits and rhizosphere soil along single roots.

Since the beginning of the industrial revolution atmospheric CO<sub>2</sub> concentrations have been steadily increasing. This probably impacts terrestrial ecosystems by stimulating plant photosynthesis and belowground allocation of the additional fixed C. Increased root exudation, promoting rhizosphere microbes, has been hypothesized as a possible explanation for the lower plant N nutritional status under elevated CO<sub>2</sub>, due to enhanced plant-microbial N competition. Legumes may counterbalance the enhanced N requirement by increased symbiotic N<sub>2</sub> fixation. The effects of elevated CO<sub>2</sub> on factors determining this symbiotic interaction were assessed in *Phaseolus vulgaris* L. grown under limited or sufficient N supply and ambient or elevated CO<sub>2</sub> concentration. Elevated CO<sub>2</sub> reduced N tissue concentrations but did not affect plant biomass production. <sup>14</sup>CO<sub>2</sub> pulse-labelling revealed no indication for a general increase in root exudation by the whole root system, which might have forced N-competition in the rhizosphere under elevated CO<sub>2</sub>. However, a CO<sub>2</sub>-induced stimulation in the exudation of sugars and malate, a chemoattractant for rhizobia, was detected in apical root zones, as potential infection sites. In nodules, elevated CO<sub>2</sub> increased the accumulation of malate as a major C source for the microsymbiont and of malonate, with functions in nodule development. Nodule biomass was also enhanced. Moreover, the release of *nod*-gene-inducing flavonoids was stimulated under elevated CO<sub>2</sub>, suggesting a selective stimulation of factors involved in establishing the *Rhizobium* symbiosis.

Since elevated-CO<sub>2</sub>-mediated effects on exudation by *Phaseolus vulgaris* L. are restricted to root apices, the abundance and function of the soil microbial community were investigated at two levels of spatial resolution to assess the response of microorganisms in the rhizosphere of the whole root system and in apical root zones to elevated CO<sub>2</sub> and different N supply. At the coarser resolution, the microbial community did not respond to CO<sub>2</sub> elevation because the C flux from the whole root system into soil did not change. At the higher spatial resolution, the



CO<sub>2</sub>-mediated enhanced root exudation from root apices led to higher enzyme activities of the C and N cycle in the adhering soil at an early stage of plant growth. At later stages, however, enzyme activities decreased under elevated CO<sub>2</sub>. This might reflect a shift in microbial C usage from the decay of polymers towards soluble carbohydrates derived from increased root exudation. CO<sub>2</sub> elevation or N supply did not affect the abundance of total and denitrifying bacteria in rhizosphere soil of apical root zones. Thus, the microbial community in the rhizosphere of bean plants responded to elevated CO<sub>2</sub> by altered enzyme regulation and not by enhanced growth.

Beyond N, plants and microorganisms may also compete for micronutrients such as Fe in the rhizosphere. *Hordeum vulgare* L., a model plant with high secretion of phytosiderophores (PS) under Fe limitation, was investigated to assess the effects of elevated CO<sub>2</sub> on PS release, Fe acquisition and potential impacts on rhizosphere microbial communities. Experiments were conducted in hydroponics and soil culture with or without Fe-fertilization and ambient or elevated CO<sub>2</sub> concentration. Elevated CO<sub>2</sub> stimulated biomass production of Fe-sufficient and Fe-deficient plants in both culture systems. Secretion of PS in apical root zones of N deficient plants increased strongly under elevated CO<sub>2</sub> in hydroponics, but no PS were detectable in root exudates from soil-grown plants. However, higher Fe shoot-contents of plants grown in soil culture without Fe supply suggest an increased efficiency for Fe acquisition under elevated CO<sub>2</sub>. Despite the evidence for altered PS secretion under elevated CO<sub>2</sub>, no significant influence on rhizosphere-bacterial communities was detected.

Low-level herbivory by parasitic nematodes is thought to induce leakage of plant metabolites from damaged roots, which can foster microorganisms. Other factors such as alterations in root exudation or morphology in undamaged roots, caused by nematode-host interactions were almost not considered yet. *Hordeum vulgare* L. was inoculated with 0, 2000, 4000 or 8000 root-knot nematodes (*Meloidogyne incognita*) for 4 weeks. In treatments with 4000 nematodes, shoot biomass, total N and P content increased by the end of the experiment. One week after inoculation, greater release of sugars, carboxylates and amino acids from apical root zones indicates leakage from this main nematode penetration site. Low levels of root herbivory stimulated root hair elongation in both infected and uninfected roots. This probably contributed to the increased sugar exudation in uninfected roots in all nematode treatments at three weeks after inoculation. Root-knots formed a separate microhabitat within the root system. They were characterised by decreased rhizodeposition and an increased fungal to bacterial ratio in the surrounding soil. This study provides evidence that, beside leakage, low-level root herbivory induces local and systemic effects on root morphology and exudation, which in turn may affect plant performance and competition.

In conclusion, this thesis extends our knowledge about the potential impact of two different plant-growth-affecting factors on rhizosphere processes, particularly at the small scale and is, thus, interesting for future assessment of management strategies in agriculture under global climate change.

## 2 Zusammenfassung

Biochemische Prozesse in der Rhizosphäre werden in hohem Maße durch organische und anorganische Verbindungen, welche passiv und/oder aktiv von Pflanzenwurzeln abgegeben werden, gesteuert. Dieser als Rhizodeposition bekannte Prozess kann durch verschiedene abiotische und biotische Faktoren stimuliert werden, was eine Anreicherung der Rhizosphäre mit leicht verfügbaren, pflanzenbürtigen Verbindungen zur Folge hat. Die vorliegende Arbeit beschäftigt sich mit den Auswirkungen der Faktoren (i) erhöhte atmosphärische CO<sub>2</sub>-Konzentration gekoppelt mit einer Limitierung verschiedener Nährstoffe im Boden und (ii) geringer Befall durch pflanzen-parasitische Nematoden auf die Menge und Zusammensetzung der Rhizodeposition. Weiterhin werden mögliche Effekte auf die pflanzliche Nährstoffaneignung und die Interaktionen zwischen Pflanze und Mikroorganismen in der Rhizosphäre näher beleuchtet. Die Arbeit umfasst vier Fall-Studien in denen Pflanzen in Wurzelkästen (Mini-Rhizotronen) kultiviert wurden. Die Verwendung dieser Kultivierungssysteme ermöglicht eine lokalisierte Sammlung von Rhizodepositen und Rhizosphärenboden entlang einzelner Wurzeln.

Seit Beginn der „Industriellen Revolution“ sind die CO<sub>2</sub>-Konzentrationen in der Atmosphäre durch menschliche Aktivitäten stetig angestiegen, was auch Auswirkungen auf terrestrische Ökosysteme haben kann. Diese umfassen unter anderem eine Stimulierung der Photosynthese und eine verstärkte Verlagerung des zusätzlich fixierten C in die Wurzeln, was zu erhöhter Wurzelexsudation führen kann. Eine verstärkte Exsudation kann mikrobielles Wachstum in der Rhizosphäre fördern und damit die Konkurrenz um Nährstoffe wie N zwischen Pflanzen und Mikroorganismen verstärken. Dies stellt eine mögliche Erklärung für den schon in früheren Studien beobachteten, geringeren N-Ernährungsstatus von Pflanzen unter erhöhten CO<sub>2</sub>-Konzentrationen dar. Leguminosen sind in der Lage, diesen erhöhten N-Bedarf durch gesteigerte, symbiotische N<sub>2</sub>-Fixierung unter „hoch-CO<sub>2</sub>“ auszugleichen. Es ist jedoch bisher unbekannt, inwieweit sich eine „CO<sub>2</sub>-Erhöhung“ auf Faktoren, welche zur Stimulierung dieser Symbiose beitragen können, auswirkt. In der ersten Studie wurden daher die Effekte von erhöhter CO<sub>2</sub>-Konzentration auf mögliche Einflussfaktoren untersucht. Als Modellpflanze wurde Bohne (*Phaseolus vulgaris* L.) ausgewählt, da sie ein gut charakterisiertes Muster von Wurzelexsudaten (Flavonoide und Carboxylate), welche Signalverbindungen bei der Induzierung der symbiotischen N<sub>2</sub>-Fixierung sind, aufweist. Die Pflanzen wurden unter zwei N-Düngungsniveaus (limitierende und ausreichende N-Zugabe) sowie unter ambienter (400 µmol mol<sup>-1</sup>) oder erhöhter (800 µmol mol<sup>-1</sup>) CO<sub>2</sub>-Konzentration kultiviert. „Hoch-CO<sub>2</sub>“ führte zu reduzierten N Spross- und Wurzelkonzentrationen, hatte aber keinen Effekt auf die

pflanzliche Biomasseproduktion. Die Ergebnisse eines  $^{14}\text{CO}_2$ -„Tracer“-Markierungsexperimentes zeigten keinen generellen Anstieg der Wurzelexsudation von der Gesamtwurzel unter erhöhter  $\text{CO}_2$ -Konzentration, welcher zu einer verstärkten Nährstoffkonkurrenz zwischen Pflanzen und Mikroorganismen in der Rhizosphäre hätte führen können. Jedoch stimulierte „hoch- $\text{CO}_2$ “ die Exsudation von Zucker und Malat, einer Locksubstanz für Rhizobien, in der apikalen Wurzelzone, den potenziellen Infektionsort für Rhizobien. Im Knöllchen-Gewebe waren unter „erhöhtem  $\text{CO}_2$ “ die Konzentrationen von Malat, die Haupt-C-Quelle des Mikro-Symbionten, und Malonat, was Funktionen bei der Knöllchenentwicklung hat, erhöht. Entsprechend war auch die Anzahl und Biomasse der Knöllchen erhöht. Darüber hinaus führte „hoch- $\text{CO}_2$ “ zu einer verstärkten Exsudation von *nod*-Gen induzierenden Flavonoiden, was ebenfalls auf eine selektive Stimulierung der Rhizobien-Symbiose hinweist.

Aufgrund der Beobachtung aus der ersten Studie, dass sich die „ $\text{CO}_2$ -Effekte“ auf die Wurzelexsudation bei Bohne auf den apikalen Wurzelraum beschränken, wurde in der zweiten Studie die Abundanz und Funktion der mikrobiellen Rhizosphärengemeinschaft auf zwei unterschiedlichen, räumlichen Ebenen untersucht. Dazu wurden die Reaktionen der Mikroorganismen auf erhöhte  $\text{CO}_2$ -Konzentration und zwei N-Düngungsstufen in der Rhizosphäre des gesamten Wurzelraumes sowie im Bereich der apikalen Wurzelzone untersucht. Die erhöhte  $\text{CO}_2$ -Konzentration hatte keinen Einfluss auf die mikrobielle Gemeinschaft in der Rhizosphäre des gesamten Wurzelraumes, was auf die unveränderte C-Exsudation der Gesamtwurzel zurückzuführen ist. Die durch „ $\text{CO}_2$ -Erhöhung“ verstärkte Exsudation in der apikalen Wurzelzone führte aber zu höheren Enzymaktivitäten des C- und N-Kreislaufes im apikalen Rhizosphärenboden. Jedoch waren zu einem späteren Entwicklungsstadium der Pflanzen diese Enzymaktivitäten unter „hoch- $\text{CO}_2$ “ erniedrigt. Dies könnte eine Verschiebung der mikrobiellen C Nutzung aus dem Abbau von Polymeren hin zur vermehrten mikrobiellen Aufnahme von leicht verfügbaren Kohlenhydraten aus verstärkter Wurzelexsudation widerspiegeln. Weder „ $\text{CO}_2$ -Erhöhung“ noch N-Zugabe hatten einen Effekt auf die Abundanz der gesamten sowie der denitrifizierenden Bakteriengemeinschaft im apikalen Rhizosphärenboden. Es lässt sich daher festhalten, dass die mikrobielle Gemeinschaft in der Rhizosphäre auf eine Erhöhung der  $\text{CO}_2$ -Konzentration mit einer Regulation von Enzymen (Produktion und Aktivität) und nicht mit verstärktem Wachstum reagierte.

Neben N, können Pflanzen und Mikroorganismen in der Rhizosphäre auch um Mikro-Nährstoffe wie Fe konkurrieren. In der dritten Studie wurde Gerste (*Hordeum vulgare* L.) als Modellpflanze, welche unter Fe-Mangel verstärkt Fe mobilisierende Phytosiderophore (PS) abgibt, ausgewählt. Ziel war es, die Auswirkungen einer erhöhten  $\text{CO}_2$ -Konzentration auf die PS-Abgabe, die pflanzliche Fe-Aneignung und mögliche Effekte auf die mikrobielle

Rhizosphärengemeinschaft zu untersuchen. Dazu wurde Gerste mit oder ohne Fe-Düngung sowie unter ambienter ( $400 \mu\text{mol mol}^{-1}$ ) oder erhöhter ( $800 \mu\text{mol mol}^{-1}$ )  $\text{CO}_2$ -Konzentration in Nährlösung oder kalkhaltigem Boden kultiviert. In beiden Kultivierungssystemen stimulierte „hoch- $\text{CO}_2$ “ die Biomasseproduktion sowohl von Pflanzen, die ausreichend mit Fe versorgt waren, als auch von Fe-Mangel Pflanzen. Außerdem wurde beobachtet, dass „erhöhtes  $\text{CO}_2$ “ zu einem starken Anstieg der PS-Abgabe in der apikalen Wurzelzone der Fe-Mangel Pflanzen aus hydroponischer Kultur führt. Es konnte aber keine PS-Abgabe der im Boden gewachsenen Pflanzen nachgewiesen werden. Jedoch zeigten die Fe-Mangel Pflanzen aus Bodenkultur höhere Fe-Gehalte im Sprossgewebe, was auf eine erhöhte Effizienz der pflanzlichen Fe-Aneignung unter „hoch- $\text{CO}_2$ “ hinweist. Trotz der deutlichen Anzeichen für eine veränderte PS-Exsudation unter erhöhter  $\text{CO}_2$ -Konzentration, konnte kein Effekt auf die bakterielle Rhizosphären-Gemeinschaft gefunden werden.

Die vierte Studie beschäftigt sich mit Herbivorie von pflanzenparasitischen Nematoden bei niedrigen Befallsraten. Frühere Studien haben gezeigt, dass das beim Eindringen der Nematoden ins Wurzelgewebe auftretende „Leakage“ pflanzenbürtiger Substanzen indirekte Auswirkungen auf die mikrobielle Gemeinschaft in der Rhizosphäre haben kann. Ob eine Interaktion zwischen Wirts-Pflanze und parasitischen Nematoden auch andere Veränderungen in Wurzelexsudation und/oder Wurzelmorphologie von nicht geschädigten Wurzeln im gleichen Wurzelsystem hervorruft, ist jedoch bisher unbekannt. In dieser Studie wurde Gerste und der „root-knot“ Nematode *Meloidogyne incognita* als Modell Pflanze-Parasit System ausgewählt. Jeweils zwei Wirtspflanzen pro Wurzelkasten wurden dabei mit 0, 2000, 4000 oder 8000 Nematoden für vier Wochen inokuliert. Durch Nematodenbehandlung (4000) wurde die Sprossbiomasse und die Spross N- und P-Gehalte der Pflanzen erhöht. Eine Woche nach Nematodenzugabe führte das „Leakage“ zu vermehrter Rhizodeposition von Zuckern, Carboxylaten und Aminosäuren in der apikalen Wurzelzone, welche den potenziellen Infektionsort für Nematoden darstellt. Die Wurzelhaarlänge im Bereich der Wurzelknötchen und auch der nicht infizierten apikalen Wurzelzonen waren unter Nematodenzugabe erhöht. Letzteres deutet auf einen systemischen, d.h. die ganze Wurzel betreffenden, Effekt des Nematodenbefalles hin, der nicht auf eine Befalls-bedingte Limitierung der N- oder P-Versorgung zurückzuführen war. Ein ähnlicher systemischer Effekt war auch drei Wochen nach Nematodenzugabe im Vergleich von nicht infizierten apikalen Wurzelzonen zu äquivalenten Wurzelzonen bei Kontrollpflanzen über eine erhöhte Zuckereksudation nachweisbar. Die Wurzelknötchen bildeten im Gesamtwurzelsystem ein eigenes Mikro-Habitat. Es wurde beobachtet, dass die Rhizodeposition der Wurzelknötchen stark reduziert war und im angrenzenden Rhizosphärenboden das Pilz zu Bakterien Verhältnis sehr stark anstieg. Diese Studie hat gezeigt, dass ein niedriger Befall der Pflanze durch „root-knot“

Nematoden, neben "Leakage", auch lokale und systemische Effekte auf Wurzelmorphologie und Exsudation hervorruft, was sich auf die Biomasseproduktion und Konkurrenzfähigkeit der Wirtspflanze auswirken kann.

Zusammenfassend lässt sich festhalten, dass die vorliegende Arbeit das Verständnis über mögliche Auswirkungen von Pflanzenwachstum beeinflussenden Faktoren auf wurzelinduzierte Mechanismen in der Rhizosphäre vertieft hat. Dieses Wissen kann zur Verbesserung von Management Strategien in der Landwirtschaft beitragen vor allem im Hinblick einer „Globalen Klimaveränderung“.

## 3 General Introduction

### 3.1 Release of organic compounds by living plant roots

Apart from their functions as organs for water and nutrient uptake as well as for anchoring the plant in soil, plant roots release a wide range of organic and inorganic compounds into their surrounding soil environment, the rhizosphere (Neumann and Römheld, 2007). In annual plant species, 20 - 60% of the photosynthetically fixed C is translocated to the roots, and a considerable proportion of this C (up to 70%) can be released into the rhizosphere. This process is termed rhizodeposition (Liljeroth, et al., 1994; Kuzyakov and Domanski, 2000). Some of this C input is rapidly metabolized by microorganisms colonising the rhizosphere (Kuzyakov, 2002). Thus, the loss of C from root epidermal and cortical cells may affect the proliferation of microorganisms within (endorhizosphere), on the surface (rhizoplane) and outside the root (ectorhizosphere) (Lynch and Whipps, 1990). Microbial concentrations in the rhizosphere can reach between  $10^{10}$  and  $10^{12}$  cells per gram soil, and invertebrate numbers in the rhizosphere are at least two orders of magnitude greater than in the surrounding soil (Lussenhop and Vogel, 1991, Brimecombe et al., 2007). Organic rhizodeposition comprises lysates, liberated by autolysis of sloughed-off cells and tissue, intact root border cells, as well as root exudates released passively (diffusates) or actively (secretions) from intact root cells (Jones et al., 2004; Neumann and Römheld, 2007). The amount of C released via root exudation ranged from 0.2 - 7% of the root dry matter per day (mean exudation value of  $150 \mu\text{g C mg}^{-1} \text{DM}$  of root growth) (Nguyen, 2003). Although root exudation probably represents a minor C transfer pathway compared to other fluxes, its biochemical nature may be more important for ecosystem functioning than its volume (Cardon, 1996). The composition of exuded compounds demonstrates a wide variety: simple and complex sugars, amino acids, organic acids, phenolics, alcohols, polypeptides and proteins, hormones and enzymes (Nguyen, 2003). The release of major low-molecular-weight organic constituents of root exudates such as sugars, amino acids, carboxylic acids and phenolics is a passive process along the steep concentration gradient, which usually exists between the cytoplasm and the external soil solution (mM vs.  $\mu\text{M}$ , respectively) (Neumann and Römheld, 2007). Besides passive diffusion, plants can respond to environmental conditions by altering their excretion of organic compounds. For example, in response to nutritional deficiency stress (e.g. P or Fe deficiency) or Al toxicity in some plant species, anion channel proteins, embedded in the plasmalemma significantly increase the passive efflux of specific carboxylates (e.g. citrate, malate, oxalate) involved in mobilizing mineral nutrients and in rhizosphere detoxification (Zhang et al., 2001; Sasaki et al., 2004; Neumann and Römheld, 2007). Together with vesicle transport, anion channels have also been

implicated in the release of Fe- and Zn-mobilizing phytosiderophores (nonproteinaceous amino acids), which are involved in the micronutrient acquisition of graminaceous plant species (Nishizawa and Mori, 1987; Sakaguchi et al., 1999). Moreover, plant roots can release a wide range of secondary metabolites such as flavonoids, which are known to play a multifunctional role in rhizospheric plant-microbe and plant-plant communications (Shaw et al., 2006). Most familiar is their function as a signal in initiating the highly specific legume-rhizobia symbiosis (Schultze and Kondorosi, 1998). In nano- to micromolar concentration, flavonoids bind to the Nod D receptor in the rhizobia-cell surface, triggering the induction of all other *nod* genes involved in producing the nodulation factors (Nod factors or lipochitooligosaccharides). Flavonoids are a diverse class of polyphenolic compounds that are produced as a result of plant secondary metabolism (Werner, 2007). Their release from roots is presumably regulated by vesicle transport and plasma membrane transporters (Walker, 2003).

Both the quality and quantity of root exudates vary considerably according to the plant species and the physiological status of the plant (Nguyen, 2003). Root exudation was observed to be highest in the phase of fast vegetative growth (Sauerbeck and Johnen, 1976). Exudation takes place along the root surface with longitudinal gradients, commonly with higher rates behind the root apices (Schilling et al., 1998; Darwent et al., 2003). This may also induce gradients in microbial degradation of root exudates (Marschner, 1995). The greater exudation at apical root zones is consistent with the concentration gradients of sugars and amino acids inside the root and with the diffusion of sugars through the apoplast from the phloem to the apical meristems (Jones, 1998, 1999). Moreover, intense root exudation, restricted to distinct root zones, may enhance the mobilization efficiency due to localized accumulation of exudate compounds in the rhizosphere. This effect may be further increased by a high capacity for nutrient uptake in apical root zones (Neumann and Römheld, 2007).

A range of multiple environmental factors influences the amount and composition of rhizodeposits. Since microbial composition and activity as well as species richness at the soil-plant interface are related either directly or indirectly to rhizodeposits, they may vary according to the same environmental factors that affect exudation. The studies presented in this thesis focus on the impact of elevated atmospheric CO<sub>2</sub> concentration accompanied by different nutrient status in soil and belowground herbivory by nematodes as factors influencing rhizosphere processes.

### **3.2 Elevated atmospheric CO<sub>2</sub> concentration**

Since 1750, anthropogenic activities, particularly burning of fossil fuels (contribution 78%) and land use changes (deforestation: 22%), have led to a marked increase (35%) in the atmospheric CO<sub>2</sub> concentration, reaching 379 μmol mol<sup>-1</sup> in 2005 (IPCC, 2007). The radiative forcing of



CO<sub>2</sub> and other greenhouse gases like methane and nitrous oxide has contributed to a 0.74°C temperature rise of the global surface since the late 19<sup>th</sup> century. Estimates of atmospheric CO<sub>2</sub> concentrations in the year 2100 range between 730 and 1088 μmol mol<sup>-1</sup> depending on economic growth, technological advances and C sequestration by biological and geological processes (IPCC, 2007). The terrestrial C cycling is probably already changing in response to these perturbations, but substantial uncertainties remain about the sensitivity of ecosystems to climate-change-forcing factors (Pendall et al., 2004a). The capacity of ecosystems to store C depends on net ecosystem production, which is the balance between net primary production and heterotrophic respiration. Increasing atmospheric CO<sub>2</sub> concentrations enhances plant C assimilation because, particularly in C<sub>3</sub> plants, the key enzyme of C assimilation, ribulose-1,5-biphosphate-carboxylase-oxygenase (RuBisCO), is unsaturated under ambient CO<sub>2</sub> conditions (Amthor et al., 2001). Thus, the rising CO<sub>2</sub> levels will have a major impact on single species and whole ecosystems (Cramer et al., 2001). Numerous studies demonstrated a stimulation of growth and biomass production of plants exposed to elevated CO<sub>2</sub> (Kimball et al., 2002; Nowak et al., 2004; Ainsworth and Long, 2005). The magnitude of the response, however, varied according to species, growing season and experimental conditions. Nutrient availability is an important factor influencing the extent of the plant response to CO<sub>2</sub> elevation. With increased plant growth, elevated CO<sub>2</sub> induces a stronger N-sink strength in plants (Zanetti et al., 1998). A reduced plant N concentration was observed for many C<sub>3</sub> species under elevated CO<sub>2</sub>. This was accompanied by an increased N-use efficiency, which was explained by an acclimation of photosynthesis to elevated CO<sub>2</sub> along with a decrease in RuBisCO and photorespiration enzymes (Webber et al., 1994; Ainsworth and Long, 2005). In C<sub>3</sub> plants about 80% of total leaf N is located in the chloroplasts, with most being bound in the RuBisCO-enzyme protein (Evans, 1989). Otherwise, a reduced plant N concentration under elevated CO<sub>2</sub> was evoked indirectly by a reduction of plant-available N in soils, reflecting intensified competition for resources between plants and soil microbes (Diaz et al., 1993). Since symbiotic N fixation could counterbalance the CO<sub>2</sub>-induced limitation of N availability, leguminous plants become favoured and may play a key role in the ecosystem response to elevated CO<sub>2</sub> (Marilley et al., 1999). The N-fixing ability of legumes is generally enhanced under elevated CO<sub>2</sub> (Lüscher et al., 2000) by two general mechanisms: formation of more nodules or higher specific N<sub>2</sub> fixation (Zanetti et al., 1998; Schortemeyer et al., 2002). Accordingly, the relative contribution of symbiotically fixed N<sub>2</sub> to total plant N increases under elevated CO<sub>2</sub> (Zanetti et al., 1996; Feng et al., 2004).

### 3.2.1 Effects of elevated CO<sub>2</sub> on belowground processes - rhizodeposition

Below-ground processes play important roles in the global C cycle because they regulate the storage of large quantities of soil organic carbon (SOC), and are potentially very sensitive to indirect effects of elevated CO<sub>2</sub>. Numerous studies have shown that increases in plant aboveground biomass under elevated CO<sub>2</sub> are often smaller than expected from the increased C fixation (Körner, 1996). It has, therefore, been hypothesized that a great part of the extra C will be allocated belowground (van Ginkel et al., 2000; Zak et al., 2000). The putative consequences for rhizodeposition, which potentially contributes to the soil C pool, are still not clear. Several studies demonstrate either an increased SOC content (Williams et al., 2000) or no influence of elevated CO<sub>2</sub> on SOC stocks even after long-term CO<sub>2</sub> enrichment (Niklaus et al., 2000; Van Kessel et al., 2000; Gill et al., 2002). In those studies, the contribution of plant-derived C (i.e. C from litter fall, rhizodeposition or root turnover) to the SOC pool was assessed collectively; thus individual fluxes were not determined. Only a few studies have investigated rhizodeposition or more specific root exudation in plant-soil systems exposed to elevated CO<sub>2</sub>, presumably due to methodological difficulties involved in quantification (Todorovic et al., 2001). With respect to total C flux from plant roots into soil, controversial results have been reported. Elevated CO<sub>2</sub> stimulated rhizodeposition in a native C3-C4 grassland ecosystem (Pendall et al., 2004b). Similarly, in a <sup>14</sup>CO<sub>2</sub> pulse-chase labelling experiment with *Lolium perenne* grown in pots with soil from the New Zealand FACE (Free Air CO<sub>2</sub> Enrichment) site, more root-derived C was found in the soil 48 h after labelling, suggesting increased root exudation (Allard et al., 2006). Enhanced root exudation is a common (Cheng and Johnson, 1998), but not a general response to elevated CO<sub>2</sub> (Hodge and Millard, 1998; Uselman et al., 2000). <sup>14</sup>CO<sub>2</sub> pulse-chase labelling of *Lolium perenne* planted into soil cores inserted within the Swiss FACE experiment revealed that C partitioning to soil was unaffected or reduced under elevated CO<sub>2</sub> (Bazot et al., 2006). These contrasting results probably reflect differences in the used plant species, the physiological state of the plant and experimental conditions (such as different N supply). For example, under non-limited N supply, C allocation to *Lolium perenne* roots was not affected by elevated atmospheric CO<sub>2</sub> (Suter et al., 2002).

An enhanced overall root exudation under elevated CO<sub>2</sub> can be caused either by stimulated root growth under elevated CO<sub>2</sub> (van Ginkel et al., 2000) or by increased exudation per unit root area (Cheng and Johnson, 1998). Qualitative alterations in the composition of root exudates towards higher C/N ratios have also been suggested under elevated CO<sub>2</sub> (Grayston et al., 1998). Moreover, elevation of CO<sub>2</sub> may improve the spatial acquisition of nutrients by stimulating fine root production (Zak et al., 2000), but may also affect the chemical availability of nutrients in the rhizosphere by increased root exudation of organic compounds, which in turn impact the

mobilization of sparingly soluble nutrients. However, this aspect has so far attracted only marginal attention. In P-deficient *Lupinus albus*, elevated CO<sub>2</sub> treatments did not affect the amount of citrate released from individual root clusters, but citrate exudation started in earlier stages of cluster root development (Wasaki et al., 2005). This was associated with a higher number of root clusters per root system, which may indicate a CO<sub>2</sub>-induced increase in the total release of P-mobilizing root exudates per plant. There was no effect on the respective rhizosphere concentrations. Most past studies have focused on the role of elevated atmospheric CO<sub>2</sub> concentration in C-, N-, and P-cycling. However, there is no information regarding CO<sub>2</sub> effects on the release rates of phytosiderophores, which are involved in the rhizosphere-availability of micronutrients, such as iron (Fe), zinc (Zn), or manganese (Mn) despite their function as essential mineral nutrients (Marschner, 1995), as factors determining pathogen resistance (Graham and Webb, 1991) and plant-microbial competition (Yehuda et al., 1996; Hördt et al., 2000).

### 3.2.2 Effects of elevated CO<sub>2</sub> on soil microbes

Soil microorganisms regulate the dynamics of organic matter decomposition and plant nutrient availability. They therefore play a key role in ecosystem response to global climate change. Whether soils can act as a sink or source for atmospheric CO<sub>2</sub> depends largely on their heterotrophic respiration of plant residues and soil organic matter (Freeman et al., 2004). Although soil microorganisms are not directly influenced by atmospheric CO<sub>2</sub> enrichment because the CO<sub>2</sub> concentrations in soil are already more than 50 times greater than in the atmosphere (van Veen et al., 1991), there may be a plant-mediated influence on microbial communities due to increased C input from rhizodeposition. The latter may enrich the rhizosphere with easily available substrates (Rogers et al., 1994). Therefore, the altered quantity and/or quality of plant-derived resources are thought to modify the activity, density and diversity of soil microbial communities. Nonetheless, the microbial responses to CO<sub>2</sub> elevation vary widely. In 47 published studies, changes in soil respiration (root and microbial respiration) under plants exposed to elevated CO<sub>2</sub> ranged from a 10% decline to a 162% increase (Zak et al., 2000). Numerous studies reported significant increases in microbial biomass, others neutral or negative responses (Zak et al., 2000; Sonneman and Wolters, 2005). The latter might be explained by limiting N availability (Diaz et al., 1993) and/or increased microbial grazing under elevated CO<sub>2</sub> (Yeates et al., 2003). Previous studies using methods targeting the overall microbial community structure by PLFA (phospholipid fatty acids) analysis or the eubacterial community diversity by 16S rDNA-based approaches (DGGE fingerprints, %G+C-base profiling) found no or weakly significant CO<sub>2</sub> effects (Griffiths et al., 1998; Hodge et al. 1998; Bruce et al., 2000; Montealegre et al., 2002; Ebersberger et al., 2004;

Wasaki et al., 2005; Drissner et al., 2007). Several authors have shown shifts in the composition of soil microbial communities, whereby those bacteria (e.g. *Pseudomonas*) colonising the rhizosphere and the rhizoplane-endorhizosphere were most affected (Schortemeyer et al., 1996; Marilley et al., 1999; Fromin et al., 2005). Moreover, elevated atmospheric CO<sub>2</sub> affects the competitive ability of root nodule symbionts (e.g. *Rhizobia*) in the legume rhizosphere, most likely leading to a selection of these particular strains to nodulate legumes (Montealegre et al., 2000; Marilley et al., 1999). Despite this, the factors involved in selectively stimulating the *Rhizobium* symbiosis at elevated CO<sub>2</sub> are not well understood. Increased root exudation of chemoattractants (malate, phenolic acids), of *nod*-gene inducing flavonoids and of other signal compounds involved in establishing the *Rhizobium* symbiosis have been discussed as possible explanations (Marilley et al., 1999).

Beyond modifying the density and structure of microbial communities, an additional input of organic substrates under elevated CO<sub>2</sub> may also affect microbial functions (Freeman et al., 2004). Extracellular enzyme activities, used to assess alterations in the function of soil microdecomposer communities (Marx et al., 2001), were sensitive indicators of CO<sub>2</sub>-induced changes in different ecosystems (Kang et al., 2001; Kandeler et al., 2006b). Elevated CO<sub>2</sub> increased the activities of enzymes such as phosphatase and urease, which promote P or N acquisition (Ebersberger et al., 2003; Drissner et al., 2007). The activity of polysaccharide-degrading enzymes also tended to increase under elevated CO<sub>2</sub> (Dhillion et al., 1996; Mayr et al., 1999). This suggests changes in soil organic matter decomposition, which may alter C sequestration and plant nutrient availability (Henry et al., 2005). With increased root exudation, much of the additional C fixed under elevated CO<sub>2</sub> is partitioned to rapidly cycling C pools rather than into more recalcitrant fractions of soil organic matter (Hungate et al., 1997). However, the addition of labile C may induce a priming effect, fostering the breakdown of more recalcitrant SOC fractions (Cheng, 1999; Hoosbeck et al., 2004).

Understanding the effects of elevated CO<sub>2</sub> on the activity of rhizosphere microorganisms is crucial because changes in nutrient turnover and particularly N cycling will feedback on plant growth. Two contrasting mechanisms concerning N-availability under elevated CO<sub>2</sub> have been hypothesized (Hu et al., 2005): (i) Diaz et al. (1993) concluded that increased C availability under elevated CO<sub>2</sub> would boost microbes, in turn immobilizing nutrients and thus constraining plant responses; (ii) Zak et al. (1993) supposed a higher N-availability for plants based on increased N mineralization rates under elevated CO<sub>2</sub> (due to priming effects of increased rhizodeposition on the mineralization of native soil organic matter). Note, however, that enhanced as well as declined gross N mineralization and N immobilization have been reported under elevated CO<sub>2</sub> (Zak et al., 2000; Richter et al., 2003; Hu et al., 2005).

The effects of CO<sub>2</sub> on denitrification processes have attracted particular attention because these are key mechanisms returning N from terrestrial ecosystems leading to a loss of plant available N in soil, while also mediating releases of the potent greenhouse gas N<sub>2</sub>O (Baggs, 2003). The complete denitrification pathway is a modular process comprising four sequential enzymatic reductions: NO<sub>3</sub><sup>-</sup> → NO<sub>2</sub><sup>-</sup> → NO → N<sub>2</sub>O → N<sub>2</sub> (see review by Philippot et al., 2007). Whereas some denitrifiers can reduce NO<sub>3</sub><sup>-</sup> into N<sub>2</sub> due to the presence of genes encoding all denitrification reductases, others have a truncated pathway (Zumft, 1997). Denitrification depends on the presence and availability of electron donors, mostly organic compounds, and of electron acceptors (nitrogen oxides, O<sub>2</sub>). All these proximate factors could be modified by plants (exudation of organic compounds, NO<sub>3</sub><sup>-</sup> assimilation) as well as by associated microbiota. Denitrifiers generally exist in soils as aerobic heterotrophs and switch to nitrate as an alternative electron acceptor only if O<sub>2</sub> is limited. This could be a competitive advantage for bacteria in the rhizosphere, where O<sub>2</sub> is limited because of root and microbial respiration (Klemedtsson et al., 1987; Højberg et al., 1996; Ghiglione et al., 2000). Moreover, an increased microbial respiration in the rhizosphere under elevated CO<sub>2</sub> may reduce the O<sub>2</sub> content and hence create anoxic sites (Zak et al., 2000). Several studies on the response of denitrification to elevated atmospheric CO<sub>2</sub> revealed increased rates of activity (Smart et al., 1997; Carnol, et al., 2002; Kettunen et al., 2005), boosting N<sub>2</sub>O emission (Baggs et al., 2003). This was explained by higher availability of root-derived C and/or by higher water use efficiency of plants under elevated CO<sub>2</sub> (Niklaus et al., 1998), which facilitate the occurrence of anaerobic conditions (Körner, 2000). We still know little about the response of microorganisms involved in denitrification to elevated CO<sub>2</sub> (Deiglmayr et al., 2004; Fromin et al., 2005; Roussel-Delif et al., 2005). Roussel-Delif et al. (2005) showed that the frequency of root-associated nitrate-dissimilating *Pseudomonas* in low fertilised *Lolium perenne* swards was increased under elevated CO<sub>2</sub>. Fromin et al. (2005) suggested that N availability may be a major factor influencing denitrifier populations under elevated CO<sub>2</sub>, most likely due to increased competition for N between plants and bacteria. Limiting NO<sub>3</sub><sup>-</sup> concentrations were also considered to be responsible for the neutral effect of elevated CO<sub>2</sub> on denitrifying processes (Martín-Olmedo et al., 2002; Barnard et al., 2004).

These variable responses of microbes to elevated CO<sub>2</sub> suggest that the complexity of the systems studied obscures a fundamental understanding of the processes involved. To date, however, no clear understanding has emerged on how root exudation (quantity and quality) is affected by elevated CO<sub>2</sub> concentrations with consequences for plant-nutrient acquisition and plant-microbial interactions in the rhizosphere.

### 3.3 Belowground herbivory by plant-parasitic nematodes

Over the last decade, the mechanisms by which plant herbivory indirectly affects soil biota and associated belowground processes like decomposition and nutrient cycling have been increasingly recognized (Bardgett et al., 1998; Van der Putten et al., 2001; Wardle et al., 2004). Aboveground herbivory can affect soil organisms and soil ecosystem-level processes significantly through the alteration of detrital inputs (e.g. plant litter) into soil but also by non-detrital inputs through changes in plant C allocation leading to altered root exudation. (Bardgett and Wardle, 2003).  $^{14}\text{C}$ - $\text{CO}_2$  pulse labelling experiments have shown that above-ground herbivory enhances root exudation (Holland et al., 1996). This in turn, may promote rhizosphere microbial populations and facilitate nutrient mineralization and plant growth (Mawdsley and Bardgett, 1997; Bardgett et al., 1998; Hamilton and Frank, 2001).

Similar processes probably also occur belowground, mediated by the root-feeding fauna (Bardgett et al., 1999a; Grayston et al., 2001). Nematodes are the most abundant and, perhaps, functionally eminent faunal group in terrestrial ecosystems. For example, their densities in grassland surface soils tend to be in the region of 3 - 4 million specimens per  $\text{m}^2$  (Yeates et al., 1997). Nematode communities in soil may also display tremendous taxonomic richness. Most of the nematode species can be allocated to specific feeding groups, in particular plant-feeders (28 - 53% of total nematodes) and bacterial-feeders (20 - 36% of total nematodes) (Bardgett et al., 1999a). Plant-parasitic nematodes are often referred to as “hidden enemies” (Cohn et al., 2002). Depending on their feeding mode, they break down root cell structure, remove cell matters, disrupt physiological processes and modify the genetic expression of their host plant (Manzanilla-López et al., 2003). While these primary effects of plant-parasitic nematodes have received considerable attention, indirect effects on rhizosphere processes are poorly understood. Yeates et al. (1998) used the  $^{14}\text{C}$ - $\text{CO}_2$  pulse labelling technique to show that, over a 14-day period, infection of *Trifolium repens* roots by low numbers of clover cyst nematodes (*Heterodera trifolii*), at average field densities (approximately 25 nematodes per plant), slightly increased the translocation of photoassimilate  $^{14}\text{C}$  to roots, and led to a significant increase in the leakage of  $^{14}\text{C}$  from roots, resulting in an enhanced microbial  $^{14}\text{C}$  content in the rhizosphere. Such infections can thus increase plant-derived C and N inputs to soil and may contribute to the regulation of microbial communities and nutrient cycling resulting in an altered availability of plant nutrients in soil. (Denton et al., 1999; Bardgett et al., 1999b; Tu et al., 2003). One suggestion is that low amounts of root infestation may reduce the overall stress on infected plants (Yeates et al., 1998; Bardgett et al., 1999a). Moreover, Dromph et al. (2006) have shown that *Heterodera trifolii* can influence vegetation characteristics by increasing the transfer of plant-derived N from their host, a N-fixing legume (*Trifolium repens*), to neighbouring grass species, thereby benefiting the growth of the neighbour.

Evidence for altered nutrient inputs to soil due to low levels of root infestation by nematodes comes mostly from grassland studies on the clover cyst nematode. Plant responses, however, may also depend on nematode feeding strategies (Yeates et al., 1999). Besides cyst nematodes including the genera *Heterodera*, the economically most important group are root-knot nematodes, particularly the genus *Meloidogyne*. The latter infect thousands of different plant species, including many crops, and may cause severe yield losses (Bird, 2004). Little, however, is known about the vigour of root-knots nematodes at low levels of root herbivory in agroecosystems. Understanding whether and how parasitic nematodes affect nutrient flow to the rhizosphere is crucial because related changes in nutrient cycling may in turn affect plant performance and competition.

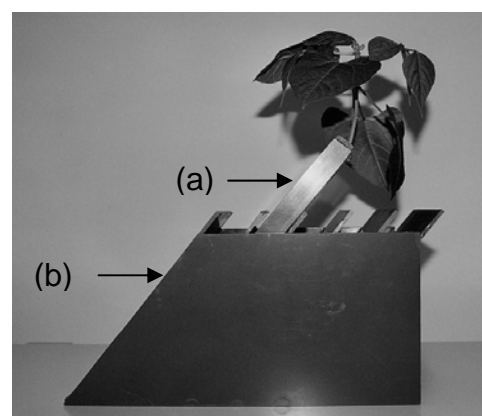
### 3.4 Collection of root exudates

Various techniques have been used to collect root exudates. Water-soluble exudates are most frequently collected from plants grown in nutrient solutions using collection techniques with trap solution. Although these methods broadly indicate exudates patterns, assigning the results to actual soil culture conditions remains difficult. For example, mechanical impedance of solid growth media as well as soil microorganisms may affect both root morphology and the total amount and composition of substances released by the root (Boeuf-Tremblay et al., 1995; Brimecombe et al., 2007; Neumann and Römheld, 2007). Alternatively, root exudates can be collected from soil-grown plants by percolating culture vessels with trap solution. Diffusion of root-derived C into soil can also be studied with two compartment systems (root mat technique), which commonly use porous membranes to separate soil-root from rhizosphere compartments (Kuzyakov, 2006). Nonetheless, these methods can not provide information on the spatial localization of the root exudation process. In many plants, root exudation is not uniformly distributed over the whole root system (Neumann and Römheld, 2007). For example, longitudinal gradients of flavonoid exudation along the roots have been observed in soybean seedlings (Kape et al., 1992), and a considerable spatial variation has been reported for the release of phytosiderophores, which predominantly takes place in apical root zones (Marschner et al., 1986). Moreover, the apical root zone – as the site of root hair formation and root elongation - is also a potential infection site for rhizobia and plant-parasitic nematodes. Microsuction cups made of HPLC capillaries were successfully used for localized collection of compounds released into the soil solution (Göttlein et al., 1996; Sandnes et al., 2005). Another approach to study spatial variation of root exudation along single roots uses sorption media with high soaking capacity (e.g. filter paper discs made of chromatography paper, nylon membranes); these are applied onto the root surface of soil-grown plants (Neumann, 2007) (Fig. 3.1). Accordingly, plants were grown in mini-rhizotrones (rhizoboxes), equipped with

removable perspex front sheets to enable access to the root surface (Fig. 3.2). The exudates can be re-extracted from the filter papers and calculation of exudation rates refers to root length (non-destructive) or biomass (destructive) of the selected root zones. The non-destructive sampling also enables repeated measurement during plant growth (Neumann, 2007). Finally, using mini-rhizotrones also enables the collection of rhizosphere soil along single roots at a small scale.



**Fig. 3.1:** Localized collection of root exudates from soil grown plants by use of sorption media (filter papers).



**Fig. 3.2:** (a) PVC mini-rhizotron (rhizobox) consisting of a corpus with irrigation holes and a perspex front lid. (b) Rhizobox holder for fixing rhizoboxes during the culture period.

A major problem of all techniques used to collect root exudates is the risk of microbial degradation during the collection period. The ability to differentiate between root exudates and microbial metabolites in the rhizosphere is also limited. Short-term collection periods (1-2 h) can minimize the risk of microbial degradation, and short-term isotopic labelling techniques (pulse-chase labelling) can help to differentiate between root exudates and microbial metabolites. Note also that only compounds dissolved in the rhizosphere solution are captured by the filter paper technique, but considerable amounts of root-derived organic compounds can be adsorbed to the soil matrix or to the cell wall. This requires recovery experiments to evaluate sorption effects to the soil matrix (Neumann, 2007). Moreover, localized sampling can also be restricted by very fine roots or low levels of soil moisture, and small sample volumes can be a limiting factor in analysing root exudates (Neumann, 2007).



## 4 Thesis Outline

This thesis uses four case studies to assess the effects of (i) elevated atmospheric CO<sub>2</sub> concentration accompanied by nutrient limitation in soil and (ii) low-level root infestation by plant parasitic nematodes on rhizodeposition and its consequences for plant-nutrient acquisition, plant-microbial interactions and nutrient cycling in the rhizosphere.

In the first study, the impact of elevated CO<sub>2</sub> on factors involved in establishing and maintaining the *Rhizobium* symbiosis in leguminous plants was examined. *Phaseolus vulgaris* L. was used as a fast-growing model plant with a well-characterized pattern of root exudates, flavonoids and carboxylates that function as signal compounds in establishing the symbiotic N<sub>2</sub> fixation. Our investigation was based on earlier studies, which revealed that many C<sub>3</sub> plants have a lower N nutritional status under elevated CO<sub>2</sub> and that leguminous plants may be able to counterbalance the enhanced N requirement by increased symbiotic N<sub>2</sub> fixation. We aimed to verify (i) whether elevated CO<sub>2</sub> limits N availability in the rhizosphere by plant-microbial competition stimulated by increased root exudation and (ii) whether in leguminous plants the lower N-nutritional status promotes symbiotic N<sub>2</sub> fixation by increased root exudation of signal compounds and/or improved carbon supply to the microsymbiont. Plants were grown under limited or sufficient N supply at ambient or elevated CO<sub>2</sub> concentrations in mini-rhizotrones. This allowed the localized collection of exudates in the apical root zone, as the potential infection site for rhizobia. Moreover, short-term <sup>14</sup>CO<sub>2</sub> pulse-labelling of photoassimilates was employed to investigate the pattern of current assimilate partitioning between plant compartments and soil C pools.

As alterations in C release from plant roots are thought to influence rhizosphere microbial growth and maintenance, the second study investigated the abundance and function of the soil microbial community. The goal was to assess the response of microorganisms to elevated CO<sub>2</sub> and different N supply in the rhizosphere of *Phaseolus vulgaris* L.. We hypothesized that specific members of the soil microbial community profit from the additional release of plant-derived C under elevated CO<sub>2</sub>. Since the first study of this thesis revealed that elevated-CO<sub>2</sub>-mediated effects on root exudation are restricted to apical root zones, this study was performed at two levels of spatial resolution. At the coarser resolution, we characterized the soil microbial community in the rhizosphere soil (total root adhering soil) using microbial biomass and basal respiration determination as well as the phospholipid fatty acids (PLFAs) pattern. At the higher spatial resolution (soil surrounding apical root zones), we focused on specific functions of the

soil microbial community involved in C, N and P cycling. Accordingly, different extracellular soil enzymes as well as densities of both total bacteria and denitrifiers were estimated.

The third experiment focused on the role of elevated atmospheric CO<sub>2</sub> concentrations on the rhizosphere-availability of micronutrients such as iron (Fe). We used *Hordeum vulgare* L. as an Fe-efficient model plant with a high potential for biosynthesis and secretion of phytosiderophores (PS) under Fe deficiency stress. The goal was to assess the effects of elevated CO<sub>2</sub> on (i) plant Fe acquisition, (ii) root-induced changes in rhizosphere chemistry and (iii) related impacts on rhizosphere microbial communities in apical root zones, which show the highest PS release rates. Experiments were conducted in hydroponic culture and in soil culture (mini-rhizotrones) with a calcareous sub-soil under ambient and elevated atmospheric CO<sub>2</sub> concentrations with and without Fe fertilization. The diversity and function of rhizosphere microbial communities were characterized by DGGE analysis of 16S rRNA and by measuring extracellular enzyme activities, respectively.

The fourth study investigated the impact of plant parasitic nematodes at low infestation rates on rhizodeposition of the host plant and related effects on rhizosphere microorganisms. Our specific objectives were to examine if a low level of root herbivory by nematodes results in (i) direct effects such as leakage from damaged roots, (ii) indirect modifications in root morphology or exudation, (iii) systemic host plant responses of undamaged root parts, and (iv) changes in the microbial community structure. The root-knot nematode *Meloidogyne incognita* and *Hordeum vulgare* L. was used as a model parasite-plant system. Plants were grown in mini-rhizotrones and were infected with different densities of *M. incognita* well below the threshold level for plant damage. Root morphology, rhizodeposition and rhizosphere microbial communities (PLFA pattern) were assessed in apical root zones, the major nematode infection site. Experiments comprised a complete life cycle of *M. incognita*, and the developed root-knots were investigated as a separate microhabitat.

**5 Elevation of atmospheric CO<sub>2</sub> and N-nutritional status modify nodulation, nodule-carbon supply, and root exudation of *Phaseolus vulgaris* L.**

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## 5.1 Abstract

Increased root exudation and a related stimulation of rhizosphere-microbial growth have been hypothesised as possible explanations for a lower nitrogen- (N-) nutritional status of plants grown under elevated atmospheric CO<sub>2</sub> concentrations, due to enhanced plant-microbial N competition in the rhizosphere. Leguminous plants may be able to counterbalance the enhanced N requirement by increased symbiotic N<sub>2</sub> fixation. Only limited information is available about the factors determining the stimulation of symbiotic N<sub>2</sub> fixation in response to elevated CO<sub>2</sub>.

In this study, short-term effects of elevated CO<sub>2</sub> on quality and quantity of root exudation, and on carbon supply to the nodules were assessed in *Phaseolus vulgaris*, grown in soil culture with limited (30 mg N kg<sup>-1</sup> soil) and sufficient N supply (200 mg N kg<sup>-1</sup> soil), at ambient (400 μmol mol<sup>-1</sup>) and elevated (800 μmol mol<sup>-1</sup>) atmospheric CO<sub>2</sub> concentrations.

Elevated CO<sub>2</sub> reduced N tissue concentrations in both N treatments, accelerated the expression of N deficiency symptoms in the N-limited variant, but did not affect plant biomass production. <sup>14</sup>CO<sub>2</sub> pulse-chase labelling revealed no indication for a general increase in root exudation with subsequent stimulation of rhizosphere microbial growth, resulting in increased N-competition in the rhizosphere at elevated CO<sub>2</sub>. However, a CO<sub>2</sub>-induced stimulation in root exudation of sugars and malate as a chemo-attractant for rhizobia was detected in 0.5-1.5 cm apical root zones as potential infection sites. Particularly in nodules, elevated CO<sub>2</sub> increased the accumulation of malate as a major carbon source for the microsymbiont and of malonate with essential functions for nodule development. Nodule number, biomass and the proportion of leghaemoglobin-producing nodules were also enhanced. The release of *nod*-gene-inducing flavonoids (genistein, daidzein and coumestrol) was stimulated under elevated CO<sub>2</sub>, independent of the N supply, and was already detectable at early stages of seedling development at 6 days after sowing.

**6 Local response of bacterial densities and enzyme activities to elevated atmospheric CO<sub>2</sub> and different N supply in the rhizosphere of *Phaseolus vulgaris* L.**

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## 6.1 Abstract

Altered flux of labile C from plant roots into soil is thought to influence growth and maintenance of microbial communities under elevated atmospheric CO<sub>2</sub> concentrations. We studied the abundance and function of the soil microbial community at two levels of spatial resolution to assess the response of microorganisms in the rhizosphere of the whole root system and of apical root zones of *Phaseolus vulgaris* L. to elevated CO<sub>2</sub> and high or low N supply.

At the coarser resolution microbial biomass C, basal respiration and phospholipid fatty acid (PLFA) patterns in the rhizosphere remained unaffected by elevated CO<sub>2</sub>, because the C flux from the whole root system into soil did not change [as shown by Haase, S., Neumann, G., Kania, A., Kuzyakov, Y., Römheld, V., Kandeler E., 2007. Elevation of atmospheric CO<sub>2</sub> and N-nutritional status modify nodulation, nodule carbon supply, and root exudation of *Phaseolus vulgaris* L. *Soil Biology & Biochemistry* 39, 2208-2221]. At a higher spatial resolution, more low molecular weight compounds were released from apical root zones under elevated CO<sub>2</sub>. Thus, at an early stage of plant growth (12 days after sowing), elevated CO<sub>2</sub> induced an increase of enzyme activities (xylosidase, cellobiosidase and leucine-aminopeptidase) in rhizosphere soil of apical root zones. At later stages of plant growth (21 days after sowing), however, enzyme activities (those above as well as N-acetyl- $\beta$ -glucosaminidase and phosphatase) decreased under elevated CO<sub>2</sub>. The abundance of total and denitrifying bacteria in rhizosphere soil of apical root zones was unaffected by CO<sub>2</sub> elevation or N supply. Plant age seemed to be the main factor influencing the density of the bacterial community. In conclusion, the soil microbial community in the apical root zone responded to elevated CO<sub>2</sub> by altered enzyme regulation (production and/or activity) and not by greater bacterial abundance.

## **7 Responses to iron limitation in *Hordeum vulgare* L. as affected by the atmospheric CO<sub>2</sub> concentration**

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## 7.1 Abstract

*Hordeum vulgare* was used as an iron-efficient model plant, with a high potential for biosynthesis and secretion of phytosiderophores (PS) under iron (Fe) deficiency stress, to assess effects of elevated atmospheric CO<sub>2</sub> concentrations on Fe acquisition, the release of PS, and potential impacts on the functional and structural diversity of rhizosphere microbial communities. Experiments were conducted in hydroponic culture and in rhizobox microcosms with a calcareous sub-soil under ambient (400 μmol mol<sup>-1</sup>) and elevated (800 μmol mol<sup>-1</sup>) atmospheric CO<sub>2</sub> concentrations with and without Fe fertilization. Elevated CO<sub>2</sub> treatments stimulated biomass production in Fe-sufficient and Fe-deficient plants, both in hydroponics and in soil culture. Root/shoot biomass ratio was increased in severely Fe-deficient plants grown in hydroponics but not under moderate Fe limitation in soil culture. Significantly increased biomass production in high CO<sub>2</sub> treatments, even under severe Fe deficiency in hydroponic culture, indicates an improved internal Fe utilization. Iron deficiency-induced secretion of PS in 0.5 - 2.5 cm sub-apical root zones was increased by 74% in response to elevated CO<sub>2</sub> treatments of barley plants in hydroponics but no PS were detectable in root exudates collected from soil-grown plants. This may be attributed to suppression of PS release by internal Fe concentrations above the critical level for Fe deficiency, determined at final harvest for soil grown barley plants, even without additional Fe supply. However, extremely low concentrations of easily plant-available Fe forms (Na-acetate extract: 0 mg kg<sup>-1</sup>, DTPA extract: 2.5 mg kg<sup>-1</sup>) in the investigated soil and low Fe seed reserves, suggest a contribution of PS-mediated Fe mobilisation from sparingly soluble Fe sources to Fe acquisition of the soil-grown barley plants during the preceding culture period. Higher Fe contents in shoots (+52%) of plants grown in soil culture without Fe supply under elevated atmospheric CO<sub>2</sub> concentrations may indicate an increased efficiency for Fe acquisition. Although, there is evidence for alterations in root growth and PS secretion rates of *Hordeum vulgare* in response to Fe deficiency and elevated atmospheric CO<sub>2</sub> concentrations, no significant influence on diversity and function of rhizosphere-bacterial communities was detectable in the outer rhizosphere soil (0 - 3 mm distance from the root surface) by DGGE of 16S rRNA gene fragments and analysis of marker enzyme activities for C-, N-, and P-cycles.



**8 Low-level herbivory by root-knot nematodes (*Meloidogyne incognita*) modifies root hair morphology and rhizodeposition in host plants (*Hordeum vulgare*)**

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## 8.1 Abstract

Low amounts of root infestation by plant parasitic nematodes are suggested to increase nutrient supply and in turn enhance microbial activity and net mineralization rate in the rhizosphere. These effects are generally related to “leakage” of plant-derived metabolites from damaged roots. Besides leakage, the present study examines other nematode-host interactions such as alterations in root exudation and morphology, which were almost not considered yet. This includes undamaged root parts in order to assess systemic plant response. The root-knot nematode *Meloidogyne incognita* (Kofoid and White, 1919; Chitwood, 1949) and barley (*Hordeum vulgare* L. cv. *Europa*) was used as model system. Host plants were grown in mini-rhizotrones inoculated with 0, 2000, 4000 or 8000 *M. incognita* for four weeks. Root morphology, rhizodeposition (sugars, carboxylates, amino acids), and rhizosphere microbial communities (PLFAs) were assessed. In treatments with 4000 nematodes, shoot biomass, total N and P content increased by the end of the experiment. Generally, an enhanced release of plant metabolites (sugars, carboxylates, amino acids) from the apical root zone occurred one week after inoculation with 4000 and 8000 *M. incognita*, indicating root leakage. Low levels of root herbivory stimulated root hair elongation in both infected and uninfected roots. These systemic changes in root morphology likely contributed to the increased sugar exudation in uninfected roots in all nematode treatments at three weeks after inoculation. Root-knots formed a separate microhabitat within the root-system. They were characterised by decreased rhizodeposition and increased fungal to bacterial ratio in the adhering rhizosphere soil. The present study provides the first evidence that, apart from leakage, nematode root herbivory at background levels induces local and systemic effects on root morphology and exudation, which in turn may affect plant performance.

## 9 General Discussion and Concluding Remarks

The present thesis investigated the effects of elevated CO<sub>2</sub> and low-level root herbivory on rhizodeposition, along with its consequences for plant-nutrient acquisition and plant-microbial interactions in the rhizosphere, based on two different model plants: *Phaseolus vulgaris* and *Hordeum vulgare*.

### 9.1 Effects of elevated atmospheric CO<sub>2</sub> concentration

This thesis yields valuable information about the influence of elevated atmospheric CO<sub>2</sub> concentration on root exudation related to nutrient acquisition by plants grown under Fe or N deficiency. Our results suggest a selective stimulation of factors involved in establishing and maintaining the *Rhizobium* symbiosis in *Phaseolus vulgaris* at elevated CO<sub>2</sub>. This was expressed in increased exudation of signal flavonoids, elevated root exudation of chemo-attractants from apical root zones (as potential infection sites for rhizobia) and increased tissue concentration of carboxylates in nodules. In *Hordeum vulgare*, elevated atmospheric CO<sub>2</sub> concentrations strongly impacted the expression of adaptations to Fe deficiency. The CO<sub>2</sub>-induced modifications comprised: increased internal Fe use efficiency, stimulated root growth and increased root exudation of Fe-mobilizing phytosiderophores from apical root zones. Both aspects – the stimulation of factors regulating symbiotic N<sub>2</sub> fixation and the improved efficiency of the Strategy II mechanism for acquiring micronutrients (Fe, Zn) in response to elevated CO<sub>2</sub> – may influence the interspecific competition in plant communities, particularly on calcareous soils with low Fe availability or on soils with low N availability. Further studies are necessary to investigate whether similar responses can also be expected under natural field conditions and in other leguminous or graminaceous plants. The case study with bean demonstrates that a CO<sub>2</sub>-induced stimulation of photosynthesis, associated with the increased shoot to root translocation of carbohydrates frequently observed in higher plants, is not related with a generally increased root exudation from the whole root system. Moreover, the CO<sub>2</sub>-mediated stimulation of root exudation in apical root zones of both model plants depends on the nutritional status and developmental stage of the plant. Therefore, future approaches for quantifying or modelling plant-derived C inputs into soil under global climate change require a detailed understanding and consideration of rhizosphere processes to obtain reliable data.

The applied localized collection techniques for root exudates and rhizosphere soil at the same spatial and temporal resolution allowed exudation from specific root zones to be linked with microbial properties in the adhering soil. Apart from the specific interactions related with the *Rhizobium* symbiosis, elevated CO<sub>2</sub> had no effects on the abundance of total and denitrifying

bacteria in the rhizosphere soil of apical root zones of bean. This contradicts our initial hypothesis that the rhizosphere microbial community profits from the additional release of plant-derived C from apical root zones. Moreover, despite the CO<sub>2</sub>-induced alterations in PS exudation of barley, subsequent experiments in soil culture revealed no effects on the structural diversity of the rhizosphere bacterial community in apical root zones. The absence of detectable CO<sub>2</sub> effects may be attributed to the marginal and/or inconsistent stimulation of root exudation due to CO<sub>2</sub> elevation during plant growth. It remains to be established whether the restriction of PS release to local short-term pulses in the apical root zones of barley (to counteract microbial degradation; Neumann and Römheld, 2007) is responsible for the limited expression of effects on rhizosphere microorganisms. In conclusion, our results revealed that future trends of increasing atmospheric CO<sub>2</sub> concentration will probably not result in major shifts or growth of rhizosphere microbial communities. Nonetheless, the study with bean clearly reveals that rhizosphere microbial communities in apical root zones may adapt to the short-term additional supply of carbohydrates by altering enzyme regulation (production and activity). This could locally modify nutrient cycling in the adhering soil.

## 9.2 Effects of low-level herbivory by plant-parasitic nematodes

Beyond the impact of elevated atmospheric CO<sub>2</sub> concentrations, the final study of this thesis indicates that biotic factors such as belowground herbivory may also affect rhizosphere processes and hence plant growth. Low-level root infestation by *Meloidogyne incognita* promoted host plant growth in barley at an infection rate of approximately 41 root knots per plant. Depending on the developmental stage of the plant-nematode interaction, low-level root infestation by *M. incognita* increased rhizodeposition of the host plant. This occurred directly via leakage from damaged plant cells or indirectly via a modified root hair morphology of uninfected root zones, which affect root exudation. These short-term local and systemic alterations in rhizodeposition potentially impact microbial communities in rhizosphere soil. In an accompanied experiment, Poll et al. (2007) observed stimulatory effects on both bacteria and fungi at an early stage of the plant-nematode interactions. Thus, low-level herbivory by *M. incognita* may affect nutrient dynamics in the rhizosphere soil and hence plant nutrient uptake, as demonstrated by increased shoot P and N contents. This in turn may affect plant performance and competition. The possibility of a beneficial effect of low-level herbivory by plant-parasitic nematodes on host plant growth is particularly tempting, considering their traditional role as pests in agricultural soils, and is thus interesting for management strategies directed at manipulating soil biota. Recent results of Mao et al. (2007) suggest that bacterial-feeding nematodes also affect root morphology through hormonal effects. The impact of such

modifications on C flow from roots to rhizosphere microbes is an interesting topic for future research in rhizosphere ecology.

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## Curriculum vitae

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09/1986 – 07/1992 Primary school in Lengefeld/Saxony  
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### Studies

10/1998 – 06/2003 Studies of Forest Science with focus on Soil Science, Site Ecology and  
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### Occupational Activity

10/2000 – 07/2001 Student assistant at the Institute of Soil Science, University of Dresden  
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- 10.-15.05.2004 „Molekulare und mikrobielle Ökologie des Bodens“, ARC Seibersdorf,  
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## Publications and Presentations

Parts of the PhD thesis and other projects were published or presented on conferences as follows:

### Poster Presentations

Haase, S., Kandeler, E., Kuzyakov, Y., Kania, A., Edelkott, I., Marschner, P., Römheld, V., Neumann, G., 2006. Localized sampling of root exudates and rhizosphere soil solution by use of sorption media – Applications and Perspectives. The First Joint Conference of the German Society for Plant Nutrition (Annual Meeting), 26.-28.09.2006, Stuttgart, Germany.

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Haase, S., Neumann, G., Kania, A., Rothe, A., Römheld, V., Kandeler, E., 2004. Root exudation and microbial diversity in the rhizosphere of bean (*Phaseolus vulgaris* L.) as affected by atmospheric CO<sub>2</sub> concentration and N-nutritional status. The Third International Nitrogen Conference (INC), 12.-16.10. 2004, Nanjing, China.

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Haase, S., Neumann, G., Kania, A., Rothe, A., Römheld, V., Kandeler, E., 2004. Effects of elevated atmospheric CO<sub>2</sub> concentration on root exudation and microbial diversity in the rhizosphere of barley and bean. Eurosoil, 04.-12.09.2004, Freiburg, Germany.

### Oral Presentations

Ruess, L., Poll, J., Marhan, S., Haase, S., Kandeler, E., 2007. Linking belowground multitrophic interactions between plants, root herbivores and microorganisms to aboveground plant performance. Annual Meeting of the Society of Ecology , 10.-14.09.2007, Marburg, Germany.

- Haase, S., Marhan, S., Neumann, G., Ruess, L., Kandeler, E., 2007. Einfluss des Nematoden *Meloidogyne incognita* auf Wurzelmorphologie, Rhizodeposition und Pflanzen-Mikroben Interaktion bei *Hordeum vulgare*. Annual Meeting of the German Society of Soil Science, 03.-7.09.2007, Dresden, Germany.
- Haase, S., 2007. Rhizodeposition und biotische Interaktionen in der Rhizosphäre von Bohne und Gerste. Colloquium of the Institute of Soil Science and Land Evaluation, 02.07.2007 University of Hohenheim, Germany.
- Haase, S., Marhan, S., Neumann, G., Ruess, L., Kandeler, E., 2006. Einfluss des Nematoden *Meloidogyne incognita* auf Wurzelmorphologie und Pflanzen-Mikroben Interaktionen bei *Hordeum vulgare*. Symposium „Ökophysiologie des Wurzelraumes“ 25.-26.09.2006, Hohenheim, Germany.
- Marhan, S., Poll, C., Haase, S., Bisharat, R., Erbs, M., Fangmeier, A., Kandeler, E., 2006. Einfluss von erhöhter atmosphärischer CO<sub>2</sub>-Konzentration auf die mikrobielle Besiedlung und den Abbau zweier Streuarten (Sommerweizen und Kornblume) in einem Ackerboden. Meeting of Commission III (Soil Biology) and Commission VIII (Soil Protection) of the German Society of Soil Science, Braunschweig, Germany.
- Haase, S., Neumann, G., Kuzyakov, Y., Römheld, V., Kandeler, E., 2005. Einfluss von erhöhter atmosphärischer CO<sub>2</sub>-Konzentration auf Wurzelexsudation und mikrobielle Diversität in der Rhizosphäre von Bohne (*Phaseolus vulgaris* L.) in Abhängigkeit der Stickstoffernährung. Annual Meeting of the German Society of Soil Science, 03.-11.09.2005, Marburg, Germany.
- Haase, S., Neumann, G., Kania, A., Rothe, A., Römheld, V., Kandeler, E., 2004. Root exudation and microbial diversity in the rhizosphere of bean (*Phaseolus vulgaris*) as affected by atmospheric CO<sub>2</sub> concentration and N-nutritional status. Symposium: Sustainable Agriculture and Land Use Strategies; 7.-8.10.2004; Peking, China
- Krumrei, S., Feger, K.H., Lorenz, K., Preston, C.M., 2003. Dynamik des Massenverlustes sowie der Elementfreisetzung bei der Zersetzung von Blatt- bzw. Nadelstreu von Kiefer, Buche, Eiche und Traubenkirsche auf einem armen Standort in der nördlichen Oberrheinebene. Annual Meeting of the German Society of Soil Science, 03.08.-07.09.2003, Frankfurt/Oder, Germany.

## Publications

### *Peer-Reviewed Journals:*

- Haase, S., Rothe, A., Kania, A., Wasaki, J., Römheld, V., Engels, V., Kandeler, E., Neumann, G., 2008. Response to iron limitation in *Hordeum vulgare* L. as affected by atmospheric CO<sub>2</sub> concentration. *Journal of Environmental Quality*, accepted.
- Haase, S., Philippot, L., Neumann, G., Marhan, S., Kandeler, E., 2008. Local response of bacterial densities and enzyme activities to elevated atmospheric CO<sub>2</sub> and different N supply in the rhizosphere of *Phaseolus vulgaris* L. *Soil Biology & Biochemistry*, in press DOI: 10.1016/j.-soilbio.2007.12.025.
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- Lorenz, K., Preston C.M., Krumrei, S., Feger, K.H., 2004. Decomposition of needle/leaf litter from Scots pine, black cherry, common oak and European beech at a conurbation forest site. *European Journal of Forest Research* 123, 177-188.
- Other Publications:*
- Kandeler, E., Haase S., Deiglmayr K., Ruess L., Marhan, S., Neumann, G., Römheld, V., Bonkowski, M., Philippot L., 2005. Small-scale distribution of soil microbial communities and enzyme processes in the rhizosphere. In: International Biomicocosmos Workshop- for the Enhancement of Rhizosphere Research. Creative Research Initiative Sousei (CRIS), Hokkaido University, Sapporo (extended abstract).
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sowie der Elementfreisetzung bei der Zersetzung von Blatt- bzw. Nadelstreu von Kiefer, Buche, Eiche und Traubenkirsche auf einem armen Standort in der nördlichen Oberrheinebene. Mitteilungen der Deutschen Bodenkundlichen Gesellschaft 103, 147-148.

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