

Chapman University

Chapman University Digital Commons

Kevin and Tam Ross Undergraduate Research
Prize

Leatherby Libraries

Spring 2022

3rd Place Contest Entry: Legume-Rhizobium Symbiosis Phenotypes

Yoobeen Lee

Teresa Hur

Isaac Min

Sydni Au Hoy

Follow this and additional works at: <https://digitalcommons.chapman.edu/undergraduateresearchprize>

Essay: Use of Leatherby Libraries's Resources

The Leatherby Libraries's extensive digital archive served to be a critical asset in the publication of our dataset, "Legume-Rhizobium symbiosis phenotypes" published on the data repository, Dryad. Over the course of three years, our goal was to assemble a dataset to investigate symbiotic phenotype studies in biological interactions between various rhizobia and legume species. Chapman University's Independent Research (291/491) provided us with a unique experience to gain competency in critical skills necessary for research and to fine-tune our ability in maneuvering the Leatherby Libraries digital archive during the COVID-19 pandemic. As current upperclassmen, we have refined our aptitude in searching for and filtering through various peer-reviewed articles to generate a dataset comprised of a constellation of rhizobia-legume focused scientific literature.

This dataset provided a unique opportunity for us to assume command of the library's arsenal of resources dedicated to conducting effective research. The instructional library session led by Dr. Doug Dechow (Science and Digital Humanities Librarian) equipped us with the necessary skill set to best utilize what the library had to offer. As freshmen, we had little to no experience with reading, much less searching for relevant journal articles. To address this, Dr. Dechow's informational session gave us insight on how to access various databases, input specific keywords, and request assistance for articles that were not readily available. We were able to take advantage of the strategies introduced to us and implement them into our database project. The session with Dr. Dechow anchored the foundation for our project and taught us the fundamental skills necessary for navigating the library website, using boolean search terms, and

leveraging its archive of databases. Furthermore, we continue to apply our knowledge not only for research but also for our additional academic endeavors.

Our project was interested in exploring scientific literature between the years of 2009 and 2020 that studied symbiotic phenotypes in the legume-rhizobium interaction. We utilized the Web of Science database, accessed through the Leathery Libraries, to generate a comprehensive list of articles that fell under our specified criteria. Reviewing documentation provided by the library such as those under the “Research and Guides” section provided insight into narrowing the search results. In particular, the “Getting Started: Library Research Strategy” documentation served as the building block for the synthesis of our dataset. It demonstrated how to effectively gather background information on a topic and deconstruct it into core key terms that could be used to generate a catalog of articles that were of interest. We searched for articles with the keywords “rhizob*,” “quantification.” and “nod*.” The ‘*’ character was included in the search terms to conduct a most exhaustive search without typing synonymous words repeatedly such as “rhizobia” and “rhizobium.” These key terms were concatenated with the “OR” function to increase our efficiency by removing the need to enter search terms in multiple places. This became the primary methodology for generating our index of scholarly articles.

The objective of our project was to synthesize an easily accessible data set that could provide relevant information on existing legume-rhizobia symbiosis research. Initially, we filtered through the articles between the years of 2009 and 2020 by reading the title and the abstract. Then, each potential article was given a thorough examination for indexing. Within the abstract, we specifically sought out mentions of plant growth-promoting rhizobacteria, rhizobia and legume species, and nodule/nodulation. After the initial round of filtering, we then assessed articles for features of interest including growth rate, density, colony, shoot dry weight, nodule

dry weight, and nodules per plant. If the article contained any of the following features, they were marked on a separate spreadsheet. Every round of filtering involved cross-checking by members of the project. This process of coding helped us conduct an accurate dataset and further expanded our knowledge of different rhizobia and legume species and reinforced key concepts.

The Leatherby Libraries and its resources have been paramount to our competency as undergraduate researchers and to the development of our dataset. Publishing this dataset has provided us with extensive experience in evaluating various scientific literature and has highlighted the significance of accessible library resources. Our time dedicated to this project has equipped us with the knowledge necessary to pursue additional endeavors in the future. We hope that this dataset will prove to be useful for fellow peers and researchers who share a deep interest in exploring the field of plant-bacterial interactions biology. The publication of the dataset has shown that the digital and physical resources that Leatherby Library provides can be fully utilized to investigate meaningful questions in diverse academic disciplines.

CSE Bibliography: Selected List

References

- Abbasi MK, Tahir MM, Azam W, Abbas Z, Rahim N. 2012. Soybean yield and chemical composition in response to phosphorus–potassium nutrition in Kashmir. *Agronomy Journal* 104:1476–1484.
- Abou-Shanab R, Wongphatcharachai M, Sheaffer C, Orf J, Sadowsky M. 2017. Competition between introduced *Bradyrhizobium japonicum* strains and indigenous bradyrhizobia in Minnesota organic farming systems. *Symbiosis* 73:155–163.
- Acosta-Jurado S, Navarro-Gómez P, Murdoch Pdel, Crespo-Rivas J-C, Jie S, Cuesta-Berrio L, Ruiz-Sainz J-E, Rodríguez-Carvajal M-Á, Vinardell J-M. 2016. Exopolysaccharide production by *sinorhizobium fredii* HH103 is repressed by Genistein in a *nodd1*-dependent manner. *PLOS ONE* 11.
- Ahmad E, Khan MS, Zaidi A. 2013. ACC deaminase producing *pseudomonas putida* strain PSE3 and *Rhizobium leguminosarum* strain RP2 in synergism improves growth, nodulation and yield of pea grown in alluvial soils. *Symbiosis* 61:93–104.
- Ahnia H, Bourebaba Y, Durán D, Boulila F, Palacios JM, Rey L, Ruiz-Argüeso T, Boulila A, Imperial J. 2018. *Bradyrhizobium Algeriense* sp. nov., a novel species isolated from effective nodules of *retama sphaerocarpa* from northeastern Algeria. *Systematic and Applied Microbiology* 41:333–339.
- Álvarez-Martínez ER, Valverde Á, Ramírez-Bahena MH, García-Fraile P, Tejedor C, Mateos PF, Santillana N, Zúñiga D, Peix A, Velázquez E. 2009. The analysis of core and symbiotic genes of rhizobia nodulating vicia from different continents reveals their common

- phylogenetic origin and suggests the distribution of *Rhizobium leguminosarum* strains together with *Vicia* seeds. *Archives of Microbiology* 191:659–668.
- Araujo J, Díaz-Alcántara C-A, Velázquez E, Urbano B, González-Andrés F. 2015. *Bradyrhizobium yuanmingense* related strains form nitrogen-fixing symbiosis with *Cajanus cajan* L. in Dominican Republic and are efficient biofertilizers to replace N fertilization. *Scientia Horticulturae* 192:421–428.
- Ardissone S, Kobayashi H, Kambara K, Rummel C, Noel KD, Walker GC, Broughton WJ, Deakin WJ. 2011. Role of *bacA* in lipopolysaccharide synthesis, peptide transport, and nodulation by *Rhizobium* sp. strain NGR234. *Journal of Bacteriology* 193:2218–2228.
- Ballhorn DJ, Younginger BS, Kautz S. 2014. An aboveground pathogen inhibits belowground rhizobia and arbuscular mycorrhizal fungi in *Phaseolus vulgaris*. *BMC Plant Biology* 14.
- Bautista VV, Monsalud RG, Yokota A. 2010. *Devosia yakushimensis* sp. nov., isolated from root nodules of *Pueraria lobata* (Willd.) Ohwi. *International Journal of Systematic and Evolutionary Microbiology* 60:627–632.
- Becerra-Rivera VA, Arteaga A, Leija A, Hernandez G, Dunn MF. 2020. Polyamines produced by *Sinorhizobium meliloti* Rm8530 contribute to symbiotically relevant phenotypes *ex planta* and to nodulation efficiency on alfalfa. *Microbiology* 166:278–287.
- Brown DB, Huang Y-C, Kannenberg EL, Sherrier DJ, Carlson RW. 2011. An *acpXL* mutant of *Rhizobium leguminosarum* bv. *Phaseoli* lacks 27-hydroxyoctacosanoic acid in its lipid A and is developmentally delayed during symbiotic infection of the determinate nodulating host *Phaseolus vulgaris*. *Journal of Bacteriology* 193:4766–4778.
- Cheng FX, Cao GQ, Wang XR, Zhao J, Yan XL, Liao H. 2009. Isolation and application of effective nitrogen fixation rhizobial strains on low-phosphorus acid soils in South China.

Science Bulletin 54:412–420.

- Chirak, ER, Kimeklis AK, Karasev ES, Kopat VV, Safronova VI, Belimov AA, Aksenova TS, Kabilov MR, Provorov NA, Andronov EE. 2019. Search for Ancestral Features in Genomes of *Rhizobium leguminosarum* bv. *viciae* Strains Isolated from the Relict Legume *Vavilovia formosa*. *Genes* 10:990.
- Cordero I, Ruiz-Díez B, Coba de la Peña T, Balaguer L, Lucas MM, Rincón A, Pueyo JJ. 2016. Rhizobial diversity, symbiotic effectiveness and structure of nodules of *Vachellia macracantha*. *Soil Biology and Biochemistry* 96:39–54.
- Costa MR, Chibeba AM, Mercante FM, Hungria M. 2018. Polyphasic characterization of Rhizobia microsymbionts of common bean [*Phaseolus vulgaris* (L.)] isolated in Mato Grosso do Sul, a hotspot of Brazilian biodiversity. *Symbiosis* 76:163–176.
- Donati AJ, Lee H-I, Leveau JH, Chang W-S. 2013. Effects of indole-3-acetic acid on the transcriptional activities and stress tolerance of *Bradyrhizobium japonicum*. *PLoS ONE* 8.
- dos Santos JMF, Alves PAC, Silva VC, Rhem MFK, James EK, Gross E. 2017. Diverse genotypes of *Bradyrhizobium nodulate* herbaceous *Chamaecrista* (Moench) (Fabaceae, Caesalpinioideae) species in Brazil. *Systematic and Applied Microbiology* 40:69–79.
- Gao M, Bengé A, Mesa JM, Javier R, Liu F-X. 2018. Use of RNA immunoprecipitation method for determining *Sinorhizobium meliloti* RNA-hfq protein associations in vivo. *Biological Procedures Online* 20.
- García Parisi PA, Lattanzi FA, Grimoldi AA, Omacini M. 2014. Multi-symbiotic systems: Functional implications of the coexistence of grass-endophyte and legume-rhizobia symbioses. *Oikos* 124:553–560.
- Goergen E, Chambers JC, Blank R. 2009. Effects of water and nitrogen availability on nitrogen

- contribution by the legume, *lupinus argenteus* pursh. *Applied Soil Ecology* 42:200–208.
- Granada CE, Beneduzi A, Lisboa BB, Turchetto-Zolet AC, Vargas LK, Passaglia LMP. 2015. Multi locus sequence analysis reveals taxonomic differences among *Bradyrhizobium* sp. symbionts of *Lupinus albescens* plants growing in arenized and non-arenized areas. *Systematic and Applied Microbiology* 38:323–329.
- Htwe AZ, Moh SM, Moe K, Yamakawa T. 2019. Biofertilizer Production for Agronomic Application and Evaluation of Its Symbiotic Effectiveness in Soybeans. *Agronomy* 9:162.
- Indrasumunar A, Menzies NW, Dart PJ. 2012. Calcium affects the competitiveness of acid-sensitive and acid-tolerant strains of *Bradyrhizobium japonicum* in nodulating and fixing nitrogen with two soybean cultivars in acid soil. *Soil Biology and Biochemistry* 46:115–122.
- Izaguirre-Mayoral ML, Sinclair TR. 2009. Irradiance regulates genotype-specific responses of rhizobium-nodulated soybean to increasing iron and two manganese concentrations in solution culture. *Journal of Plant Physiology* 166:807–818.
- Kajic S, Krznanic Dora, Rajnovic I, Sikora S. 2020. Isolation and characterization of endophytic bacteria from soybean (*Glycine max* L.). *Journal of Central European Agriculture* 21:151-158.
- Kalita M, Malek W. 2017. Molecular phylogeny of *Bradyrhizobium* bacteria isolated from root nodules of tribe Genisteeae plants growing in southeast Poland. *Systematic and Applied Microbiology* 40:482–491.
- Koch M, Delmotte N, Ahrens CH, Omasits U, Schneider K, Danza F, Padhi B, Murset V,

- Braissant O, Vorholt JA, et al. 2014. A link between Arabinose utilization and Oxalotrophy in bradyrhizobium japonicum. Applied and Environmental Microbiology 80:2094–2101.
- Li H, Deng Y, Wu T, Subramanian S, Yu O. 2010. Misexpression of MIR482, Mir1512, and MIR1515 increases soybean nodulation . Plant Physiology 153:1759–1770.
- Li YZ, Wang D, Feng XY, Jiao J, Chen WX, Tian CF. 2016. Genetic analysis reveals the essential role of nitrogen phosphotransferase system components in sinorhizobium fredii CCBAU 45436 symbioses with soybean and Pigeonpea Plants. Applied and Environmental Microbiology 82:1305–1315.
- Liu LX, Li QQ, Zhang YZ, Hu Y, Jiao J, Guo HJ, Zhang, XX, Zhang B, Chen WX, Tian CF. 2017. The nitrate-reduction gene cluster components exert lineage-dependent contributions to optimization of *Sinorhizobium* symbiosis with soybeans. Environmental Microbiology 19:4926–4938.
- Martín-Rodríguez JÁ, Leija A, Formey D, Hernández G. 2018. The microrna319d/TCP10 node regulates the common bean – rhizobia nitrogen-fixing symbiosis. Frontiers in Plant Science 9.
- Martínez-Salazar JM, Sandoval-Calderón M, Guo X, Castillo-Ramírez S, Reyes A, Loza MG, Rivera J, Alvarado-Affantranger X, Sánchez F, González V, et al. 2009. The rhizobium etli RPOH1 and RPOH2 sigma factors are involved in different stress responses. Microbiology 155:386–397.
- Mohammed MA, Chernet MT, Tuji FA. 2020. Phenotypic, stress tolerance, and plant growth promoting characteristics of rhizobial isolates of grass pea. International Microbiology 23:607–618.

- Muñoz V, Ibáñez F, Tordable M, Megías M, Fabra A. 2014. Role of reactive oxygen species generation and nod factors during the early symbiotic interaction between bradyrhizobia and peanut, a legume infected by crack entry. *Journal of Applied Microbiology* 118:182–192.
- Naluyange V, Ochieno DMW, Maingi JM, Ombori O, Mukaminega D, Amoding A, Odendo M, Okoth SA, Shivoga WA, Muoma JVO. 2014. Compatibility of rhizobium inoculant and water hyacinth compost formulations in Rosecoco Bean and consequences on *Aphis Fabae* and *Colletotrichum lindemuthianum* infestations. *Applied Soil Ecology* 76:68–77.
- Nascimento FX, Brígido C, Glick BR, Oliveira S, Alho L. 2012. *Mesorhizobium Ciceri* LMS-1 expressing an exogenous 1-aminocyclopropane-1-carboxylate (ACC) deaminase increases its nodulation abilities and chickpea plant resistance to soil constraints. *Letters in Applied Microbiology* 55:15–21.
- Nguyen TH, Brechenmacher L, Aldrich JT, Clauss TR, Gritsenko MA, Hixson KK, Libault M, Tanaka K, Yang F, Yao Q, et al. 2012. Quantitative phosphoproteomic analysis of soybean root hairs inoculated with *bradyrhizobium japonicum*. *Molecular & Cellular Proteomics* 11:1140–1155.
- Novák K, Lisá L, Škrdleta V. 2010. Pleiotropy of PEA RISFIXC supernodulation mutation is symbiosis-independent. *Plant and Soil* 342:173–182.
- Otsubo AA, Brito OR, Mercante FM. 2013. Productivity and nodulation of promising lineages of the Carioca Bean Group inoculated with *Rhizobium Tropici* or supplemented with nitrogen fertilizer. *Semina: Ciências Agrárias* 34:2763.

- Qiao Z, Zogli P, Libault M. 2019. Plant Hormones Differentially Control the Sub-Cellular Localization of Plasma Membrane Microdomains during the Early Stage of Soybean Nodulation. *Genes* 10:1012
- Quelas JI, Mongiardini EJ, Casabuono A, López-García SL, Althabegoiti MJ, Covelli JM, Pérez-Giménez J, Couto A, Lodeiro AR. 2010. Lack of galactose or galacturonic acid in *bradyrhizobium japonicum* USDA 110 exopolysaccharide leads to different symbiotic responses in soybean. *Molecular Plant-Microbe Interactions*® 23:1592–1604.
- Ramirez-Bahena, MH, Flores-Felix, JD, Velazquez E, Peix A. 2020. The Mimosoid tree *Leucaena leucocephala* can be nodulated by the symbiovar *genistearum* of *Bradyrhizobium canariense*. *Systematic and Applied Microbiology* 43:126041.
- Suri VK, Choudhary AK. 2013. Effects of vesicular arbuscular mycorrhizae and applied phosphorus through targeted yield precision model on root morphology, productivity, and nutrient dynamics in soybean in an acid alfisol. *Communications in Soil Science and Plant Analysis* 44:2587–2604.
- Sánchez C, Gates AJ, Meakin GE, Uchiumi T, Girard L, Richardson DJ, Bedmar EJ, Delgado MJ. 2010. Production of nitric oxide and nitrosylhemoglobin complexes in soybean nodules in response to flooding. *Molecular Plant-Microbe Interactions*® 23:702–711.
- Sánchez-Pardo B, Zornoza P. 2014. Mitigation of CU stress by legume–rhizobium symbiosis in white lupin and soybean plants. *Ecotoxicology and Environmental Safety* 102:1–5.
- Van Noorden G, Verbeek R, Dinh Q, Jin J, Green A, Ng J, Mathesius U. 2016. Molecular signals controlling the inhibition of nodulation by nitrate in *medicago truncatula*. *International Journal of Molecular Sciences* 17:1060.
- Vankosky MA, Cárcamo HA, Dossdall LM. 2011. Response of *Pisum sativum* (fabales:

- Fabaceae) to *Sitona lineatus* (Coleoptera: Curculionidae) infestation: Effect of adult weevil density on damage, larval population, and yield loss. *Journal of Economic Entomology* 104:1550–1560.
- Vargas-Diaz AA, Ferrera-Cerrato R, Silva-Rojas HV, Alarcon A. 2019. Isolation and evaluation of endophytic bacteria from root nodules of *Glycine max* L. (Merr.) and their potential use as biofertilizers. *Spanish Journal of Agricultural Research* 17:e1103.
- Wang YC, Wang F, Hou BC, Wang ET, Chen WF, Sui XH, Chen WX, Li Y, Zhang YB. 2013. Proposal of *Ensifer Psoraleae* sp. nov., *Ensifer Sesbaniae* sp. nov., *Ensifer Morelense* Comb. nov. and *Ensifer Americanum* comb. Nov. *Systematic and Applied Microbiology* 36:467–473.
- Wendlandt CE, Regus JU, Gano-Cohen KA, Hollowell AC, Quides KW, Lyu JY, Adinata ES, Sachs JL. 2019. Host investment into symbiosis varies among genotypes of the legume *Acmispon strigosus*, but host sanctions are uniform. *New Phytologist* 221:446–458.
- Yuan SL, Li R, Chen HF, Zhang CJ, Chen LM, Hao QN, Chen SL, Shan ZH, Yang ZL, Zhang XJ, Qiu DZ, Zhou XA. 2017. RNA-Seq analysis of nodule development at five different developmental stages of soybean (*Glycine max*) inoculated with *Bradyrhizobium japonicum* strain 113-2. *Scientific Reports* 7:42248.
- Yuan T, Liu L, Huang S, Taher AH, Tan Z, Wu G, Peng G. 2018. *Rhizobium Wuzhouense* sp. nov., isolated from roots of *Oryza officinalis*. *International Journal of Systematic and Evolutionary Microbiology* 68:2918–2923.
- Zepeda I, Sánchez-López R, Kunkel JG, Bañuelos LA, Hernández-Barrera A, Sánchez F, Quinto C, Cárdenas L. 2014. Visualization of highly dynamic F-actin plus ends in growing *Phaseolus vulgaris* root hair cells and their responses to *Rhizobium Etli* Nod factors. *Plant*

- and Cell Physiology 55:580–592.
- Zhang X, Wang L, Li J, Batstone RI, Frederickson ME. 2020. *Medicago truncatula* adjusts root proliferation, nodule formation, and partner choice in response to local N heterogeneity. *Plant and Soil* 450:417–428.
- Zhang YM, Li Y, Chen WF, Wang ET, Sui XH, Li QQ, Zhang YZ, Zhou YG, Chen WX. 2012. *Bradyrhizobium Huanghuaihaiense* sp. nov., an effective symbiotic bacterium isolated from soybean (*Glycine Max* L.) nodules. *International Journal of Systematic and Evolutionary Microbiology* 62:1951–1957.
- Zhang YZ, Wang ET, Li M, Li QQ, Zhang YM, Zhao SJ, Jia XL, Zhang LH, Chen WF, Chen WX. 2011. Effects of rhizobial inoculation, cropping systems and growth stages on endophytic bacterial community of Soybean Roots. *Plant and Soil* 347:147–161.
- Zhao CZ, Huang J, Gyaneshwar P, Zhao D. 2018. *Rhizobium* sp IRBG74 Alters *Arabidopsis* Root Development by Affecting Auxin Signaling. *Frontiers in Microbiology* 8.
- Zhao H, Chang QS, Zhang DX, Fang RJ, Zhao H, Wu FY, Wang XM, Lu GH, Qi JL, Yang YH. 2015. Overexpression of *lemyb1* enhances Shikonin Formation by up-regulating key shikonin biosynthesis-related genes in *Lithospermum Erythrorhizon*. *Biologia plantarum* 59:429–435.
- Zhou PF, Chen WM, Wei GH. 2010. *Mesorhizobium Robiniae* sp. nov., isolated from root nodules of *Robinia pseudoacacia*. *International Journal of Systematic and Evolutionary Microbiology* 60:2552–2556.
- Zimmer S, Messmer M, Haase T, Piepho H-P, Mindermann A, Schulz H, Habekuß A, Ordon F, Wilbois K-P, Heß J. 2016. Effects of soybean variety and *Bradyrhizobium* strains on yield, protein content and biological nitrogen fixation under cool growing conditions in

Germany. *European Journal of Agronomy* 72:38–46.

Summary of Research Project: Legume-Rhizobium symbiosis phenotypes

Our project aimed to examine scientific literature and produce an accurate dataset reflecting the various legume-rhizobium phenotypes observed in prior research. We coded articles published between January 1, 2009, and September 30, 2020, based on the presence of legume or rhizobium phenotypes. Our focus concentrated on papers that quantified rhizobia and identified relevant legume phenotypes. We used the Web of Science database through Chapman University's Leatherby Libraries to search for legume-rhizobium symbiosis articles to access necessary literature. Within the database, keywords Topic = (rhiz* OR bradyrhiz* OR mesorhiz* OR sinorhiz* OR ensifer*) AND Topic = (count* OR form* OR quant* OR estimat*) AND Topic = nod* were searched in the engine for desired articles. Initially yielding 2,440 papers, the author, title, source, year of publication, and DOI were exported via Web of Science. Then, the filtering of journal articles began by solely examining the titles and abstracts. Articles that included mentions of plant growth-promoting rhizobacteria, rhizobia and legume species, and nodule/nodulation within the abstract were downloaded for coding. Subsequently, we read through the literature to identify 25 different legume-rhizobium phenotypes including: "Rhizobia Growth or Growth Rate," "Plant Growth or Dry Weight or Fresh Weight," "Root Dry Weight or Density," "Root Branching or Length," Rhizobia Colony Morphology," "Rhizobia Number," "Shoot Dry Weight," "Nodule Dry Weight," "Number of Nodules Per Plant," "Number of Nodules," "Acetylene Reduction," "Isotopic 15N," "Biomass," "Chlorophyll," "Roots Per Plant," "Leaf Number or Weight," "Shoot Length," "Nitrogen Fixation," "Nodule Occupancy or Diversity," "Seed Count or Production," "Nodule Structure," "Shoot Number," "Root/Shoot Ratio," "Branch Dry Weight," and "Nodule Color." In total, 771 articles were analyzed for relevant phenotypes.