

Texas State Soil and Water Conservation Board State General Revenue Nonpoint Source Grant Program FY2013 Project Workplan 13-50

| | PROJECT | SUMMARY PAGE | | | |
|--|---|--|---|--|--|
| Title of Project | Statewide Bacterial Sourc | e Tracking Program for FYs 2013-2014 | | | |
| Project Goals | Support BST analyses across the State through (1) maintenance of analytical infrastructure at public BST laboratories; (2) continued development and implementation of statewide BST template-SOPs; (3) delivery of informational materials on bacteria BMPs and the use and applicability of BST and the State-supported analytical labs; (4) further expansion and evaluation of the Texas <i>E. coli</i> BST Library; and (5) further development of suitable source-specific bacterial markers for library independent BST. | | | | |
| Project Tasks | (1) Project Administration Collection; (4) Analytical Development; (5) Outreac | n; (2) Quality Assurance; (3) Known Source Laboratory Capacity, Library Expansion, a ch on Bacterial Source Tracking and BMPs | e Fecal Sample and Methods | | |
| Measures of Success | Updated BST template-SOPs for ERIC-RP, RP and <i>Bacteroidales</i> PCR Data analyzed for approximately 100 known source <i>E. coli</i> isolates from TSSWCB project 11-50 for expansion of the Texas <i>E. coli</i> BST Library Fingerprinting and analysis of 20 known-source <i>E. coli</i> isolates collected as part of TSSWCB project 11-51 Expansion and evaluation of the Texas <i>E. coli</i> BST Library through analysis of approximately 100 targeted known source fecal samples Evaluation of geographical and temporal stability of the Texas <i>E. coli</i> BST Library and diversity of source specific isolates Development/evaluation of source-specific bacterial markers for library-independent BST Outreach through website and delivery of BST and BMP informational materials to | | | | |
| Project Type | Implementation (); Educa | ation (); Planning (); Assessment (X); Gro | undwater () | | |
| Status of Waterbody on 2010 Texas Integrated Report | Segment ID Statewide | Parameter of Impairment or Concern bacteria | Category 4 and 5 | | |
| Project Location (Statewide or Watershed and County) | Statewide | | | | |
| Key Project Activities | Hire Staff (X); Surface W Education (); Implementa Demonstration (); Plannin | <pre>fater Quality Monitoring (); Technical Assi ation (); BMP Effectiveness Monitoring () ng (); Modeling (); Bacterial Source Track</pre> | stance (); ; sing (X); Other () | | |
| 2012 Texas NPS Management Program Reference | Component 1 – LTG Objectives 1, 2, 3, 6 Component 1 – STG 1C Components 2, 3, 5 | | | | |
| Project Costs | \$415,348 | | | | |
| Project Management | Texas Water Resource The University of Texas Regional Car Texas A&M AgriLife Texas A&M Institute | ces Institute xas Health Science Center at Houston Scho mpus e Research, Department of Soil and Crop S of Renewable Natural Resources | ool of Public Health, ciences | | |
| Project Period | October 1, 2012 – Septem | lber 30, 2014 | | | |

Part I – Applicant Information

Applicant

| Project Lea | .d | Kevin Wagner, Ph | .D. | | | | | | |
|-------------------------------|---------|--|--------------|------|--------|-------|------------|--------|--|
| Title | | Associate Director | | | | | | | |
| Organizatio | on | Texas Water Resou | arces Instit | tute | | | | | |
| E-mail Add | lress | klwagner@ag.tamu | 1.edu | | | | | | |
| Street Add | ess | 1500 Research Pky | vy, Ste A2 | 40 | | | | | |
| | | 2118 TAMU | | | | | | | |
| City | College | Station County Brazos State TX Zip Code 77843-2118 | | | | | 77843-2118 | | |
| Telephone Number 979-845-2649 | | | | | Fax Nu | umber | 979-84 | 5-8554 | |

| Co-Applic | ant | | | | | | | | |
|------------------|---------|---|-----------|---------|--------|-------|-------------|----------|-------|
| | | | | | | | | | |
| Project Lea | d | George Di Giovani | ni, Ph.D. | | | | | | |
| Title | | Professor | | | | | | | |
| Organizatio | on | The University of Texas Health Science Center at Houston School of Public Health, El Paso | | | | | th, El Paso | | |
| | | Regional Campus | | | | | | | |
| E-mail Add | lress | george.d.digiovanr | ni@uth.tm | c.edu | | | | | |
| Street Add | ess | 1101 N. Campbell, CH 412 | | | | | | | |
| City | El Paso | | County | El Paso | State | TX | | Zip Code | 79902 |
| Telephone | Number | 915-747-8509 | | | Fax Nu | umber | 915-74 | 7-8512 | |

| Co-Applic | ant | | | | | | | | |
|---|---------|--|-------------|-------------|-----------|--------------|----------|----|--|
| Project Lea | ıd | Terry Gentry, Ph.I |) . | | | | | | |
| Title | | Assistant Professor | • | | | | | | |
| Organizatio | on | Texas A&M AgriL | life Resear | ch, Departn | nent of S | oil and Crop | o Scienc | es | |
| E-mail Add | lress | tgentry@ag.tamu.e | edu | | | | | | |
| Street Add | ess | 370 Olsen Blvd | | | | | | | |
| | | 2474 TAMU | | | | | | | |
| City | College | Station County Brazos State TX Zip Code 77843-2474 | | | | | | | |
| Telephone Number 979-845-3041 Fax Number 979-845-0456 | | | | | | | | | |

| Co-Applic | ant | | | | | | | | |
|------------------|---------|--|----------|-------|--------|-------|----------|----------|-------|
| | | | | | | | | | |
| Project Lea | d | Roel Lopez, Ph.D. | | | | | | | |
| Title | | Interim Director | | | | | | | |
| Organizatio | on | Texas A&M Institute of Renewable Natural Resources / Texas Water Resources Institute | | | | | nstitute | | |
| E-mail Add | lress | roel@tamu.edu | | | | | | | |
| Street Add | ess | 4040 Broadway, St | uite 360 | | | | | | |
| City | San Ant | onio | County | Bexar | State | TX | | Zip Code | 78209 |
| Telephone | Number | 210-277-0292, et | xt. 202 | | Fax Nu | umber | 210-27 | 7-0378 | |

| Names | Roles & Responsibilities |
|---|--|
| Texas State Soil and Water Conservation | Provide state oversight and management of all project activities and |
| Board (TSSWCB) | ensure coordination of activities with related projects. |
| Texas Water Resources Institute (TWRI) | Project Coordination and Administration, Project Reporting, and |
| | Outreach (Tasks 1 and 5). |
| The University of Texas Health Science Center | Work in conjunction with AgriLife SCSC to perform all work |
| at Houston School of Public Health, El Paso | described in Tasks 2, 3 and 4. |
| Regional Campus (UTSPH EP) | |
| Texas A&M AgriLife Research – Department | Work in conjunction with UTSPH EP to perform all work described |
| of Soil and Crop Sciences (AgriLife SCSC) | in Tasks 2, 3 and 4. |
| Texas A&M Institute of Renewable Natural | Work in conjunction with UTSPH EP and AgriLife SCSC to perform |
| Resources (IRNR) | work described in Task 3. |

Part II – Project Information

| Watershed Information | | | | |
|------------------------------|------------------------------------|------------|---------------------|--------------|
| Watershed or Aquifer Name(s) | Hydrologic Unit Code (12 Digit) | Segment ID | Category on 2010 IR | Size (Acres) |
| Statewide | N/A | N/A | 4 and 5 | N/A |

Water Quality Impairment

Describe all known causes (i.e., pollutants of concern) and sources (e.g., agricultural, silvicultural) of water quality impairments or concerns from any of the following sources: *2010 Texas Integrated Report*, Clean Rivers Program Basin Summary/Highlights Reports, or other documented sources.

The 2010 303(d) List identified >300 contact recreation use impairments (waterbody-pollutant combinations) and 15 oyster water use impairments due to excessive bacteria (*E. coli, Enterococcus spp.*, or fecal coliform). These bacteria impairments account for more than half of all impairments on the 2010 303(d) List. This is more than 3 times as many impairments as the next largest number of a specific impairment type/pollutant. These indicator bacteria originate from human (WWTF, OSSF) and animal (wildlife, pets, livestock, feral hogs) sources and reach waterbodies through point source discharges, direct deposition, and NPS runoff.

Project Narrative

Problem/Need Statement

Protection of water resources is one of the most significant environmental challenges of the new millennium. Nonpoint sources (NPS) of pollution, including agricultural activities, can greatly impact water quality. One key component in effectively implementing a NPS pollution abatement program is the identification and assessment of sources of fecal pollution. Proper evaluation of these sources is needed to target best management practices (BMPs) and develop bacterial total maximum daily loads (TMDLs) or watershed protection plans (WPPs). This information may also be useful to properly assess risk in contact recreation, as many waterborne pathogens causing human illness do not colonize nonhuman hosts. According to the *2010 Texas Integrated Report*, there are over 300 impairments due to excessive bacteria.

Fecal coliform bacteria have extensively been used as an indicator of fecal pollution and the potential presence of other pathogenic microorganisms in water. It has been established that the fecal coliform bacterium *E. coli* is more closely associated with fecal pollution than other fecal coliform bacteria, which may normally reside and multiply in the environment. *E. coli* is a common inhabitant of animal and human intestines and recent studies have shown that isolates from humans and various host animals (e.g., cattle, chickens, and pigs) may differ genetically and phenotypically. Use of genetic and biochemical tests may allow the original host species to be identified and is referred to as bacterial source tracking (BST).

The premise behind BST is that genetic and phenotypic tests can identify bacterial strains that are host specific so that the original host species and source of the fecal contamination can be identified. Often *E. coli* or *Enterococcus* spp. are used as the bacteria targets in BST, as this provides a direct link with water quality standards which are usually based on one of these two indicators (Parveen, Portier et al. 1999; Dombek, Johnson et al. 2000; Graves, Hagedorn et al. 2002; Field, Chern et al. 2003; Hartel, Summer et al. 2003; Kuntz, Hartel et al. 2003; Stoeckel, Mathes et al. 2004; Harwood, Levine et al. 2005). While there has been some controversy concerning host specificity and survival of *E. coli* in the environment (Gordon, Bauer et al. 2002), this indicator organism has the advantage that it is known to correlate with the presence of fecal contamination and is used for human health risk assessments. BST of *E. coli*, therefore, has the advantages of direct regulatory significance and availability of standardized culturing techniques for water samples, such as EPA Method 1603 (EPA 2005).

BST is a valuable tool for identifying human and animal sources of fecal pollution. Comprehensive BST has been completed by UTSPH EP (formerly with Texas A&M AgriLife Research) for (1) the Lake Waco and Belton Lake watersheds, (2) several San Antonio area watersheds, (3) the Lake Granbury watershed, (4) Buck Creek, and (5) the Leon and Lampasas Rivers watersheds. The Waco/Belton and Buck Creek studies were funded by the TSSWCB through Clean Water Act §319(h) NPS grants from the U.S. Environmental Protection Agency (EPA) (TSSWCB projects 02-10 and 06-11, respectively) and the Leon and Lampasas project through state general revenue funds (TSSWCB project 10-51); while the San Antonio study and Lake Granbury studies were funded by the Texas Commission on Environmental Quality (TCEQ). In addition, AgriLife SCSC has completed BST projects for the Little Brazos River tributaries and Big Cypress Creek watersheds (TSSWCB projects 09-52 and 09-55, respectively). Additionally, with TSSWCB funding, BST projects are currently under way in the Leona River and Attoyac Bayou watersheds to assess water quality impairments (projects 11-50 and 09-10, respectively). A Texas E. coli BST Library has been developed based on known source isolates from the Waco/Belton, San Antonio, Granbury, Buck Creek, Big Cypress, Little Brazos River, Attoyac Bayou, Leon River, Lampasas River, Upper Trinity River and Upper Oyster Creek watersheds. The Texas E. coli BST Library (ver. 8-12) currently contains 1,669 E. coli isolates obtained from 1,455 different domestic sewage, wildlife, livestock and pet fecal samples. While this represents a significant step towards development of a statewide E. coli BST library, continued expansion of the library to include additional known source isolates from different Texas watersheds and different animal hosts is still needed. This will allow continued evaluation of the library for geographical stability and the diversity of source specific isolates to identify specific needs for future expansion and refinement of the library. The use of the Texas E. coli BST Library will provide for significant cost and time savings for the identification of NPS pollution in the development of TMDLs and WPPs.

A Task Force on Bacteria TMDLs was jointly established by the TSSWCB and the TCEQ in fall 2006. In the Task Force's Report, a strategy to address current and future bacterial TMDLs and Implementation Plans (I-Plans) was outlined. The Task Force describes and makes recommendations for effective use of BST methods that have been used in Texas. These include enterobacterial repetitive intergenic consensus sequence polymerase chain reaction (ERIC-PCR), RiboPrinting (RP), Kirby-Bauer antibiotic resistance analysis (KB-ARA), carbon source utilization (CSU), and *Bacteroidales* PCR. The Task Force recommended using library-independent methods such as *Bacteroidales* PCR for preliminary qualitative analyses and library-dependent methods (e.g., ERIC-PCR and RP) if more quantitative data are required. Further characterization of known source *E. coli* for expansion of the Texas *E. coli* BST Library and continued support of established BST analytical infrastructure will help achieve the recommendations of the Task Force.

The Task Force Report identified certain Research and Development (R&D) needs to advance understanding of bacteria. Specifically, 30 types of studies or research needs in 6 categories (including Characterization of Sources and Bacterial Source Tracking) were identified. This list was not exhaustive and no attempt was made to prioritize these activities. As such, there is a need to update, expand and prioritize these BST-related R&D activities.

Lastly, the state of BST science, methodologies, application and confidence has evolved greatly in the past few years. A host of new information is currently available, yet not readily distributed or known to state and federal agency personnel. To address this, the *2012 BST – State of the Science Conference* was held. To build on the success of this conference, continued outreach and technology transfer is needed to foster dialogue and collaboration and bring water resource managers up to speed on advances in BST technologies, methodologies, applications and results.

Project Narrative

General Project Description (Include Project Location Map)

The Texas *E. coli* BST Library is dynamic, with new isolates being added with each successive BST project. The current library (ver. 8-12) contains known source isolates from over a dozen watersheds, as well as wildlife isolates from South Texas. Under this project, ERIC-RP data for approximately 100 known source *E. coli* isolates from the Leona River watershed (TSSWCB Project 11-50) will be provided by AgriLife SCSC to UTSPH EP for analysis and expansion of the state library. In addition, approximately 100 known source fecal samples from targeted animal sources will be collected and analyzed for *E. coli* to further expand the state library and provide additional datasets for library evaluation. In particular, the state library has very few *E. coli* from wildlife species such as mice, rabbits, nutria and squirrels. By collecting some of these known source samples from a previously studied watershed (e.g., Leon River watershed), the temporal stability of the library will also be assessed. The geographic stability of the library will be evaluated by performing watershed exclusive and inclusive statistical analyses. In addition, the fingerprint diversity of source-specific *E. coli* isolates will be investigated to help evaluate the strain representativeness of the library. This will allow the project team to identify specific needs for the future expansion and refinement of the library.

There have been significant developments in library-independent BST methods, including bacterial genetic markers specific to different animal sources and humans (i.e. Bernhard and Field 2000; Dick, Bernhard et al. 2005; Scott, Jenkins et al. 2005; Hamilton, Yan et al. 2006). Library-independent methods are cost-effective, rapid, and potentially more specific than library-dependent methods. Concerns with many of the recently developed library-independent approaches include uncertainties regarding geographical stability of markers and the difficulty of interpreting results in relation to regulatory water quality standards and microbial risk, since some target microorganisms are not regulated. More importantly, these library-independent methods can only detect a limited range of pollution sources and are currently only semi-quantitative. For example, the Bacteroidales PCR (Bernhard and Field 2000; Dick, Bernhard et al. 2005) can detect fecal pollution from ruminants, humans, dogs, horses and pigs; but currently no further discrimination is possible. Despite these limitations, this method may be very useful for the rapid and inexpensive assessment of the possible sources of fecal pollution impacting a waterbody. UTSPH EP (under TSSWCB project 10-50) has generated promising preliminary results for a *Bacteroidales* PCR method to detect feral hog fecal pollution, as well as identified possible genetic targets for discriminating human and animal E. coli. A simple library-independent method for distinguishing human from animal E. coli would be quite useful for BST studies. Current research in this area at UTSPH EP is based on sequence analysis of ERIC-PCR products from isolates identified through data mining of the Texas E. coli BST Library. Library-independent source-specific methods have recently been described for poultry (Weidhaas et al. 2010) and cattle (Shanks et al. 2010). Importantly, UTSPH EP has observed some cross-reactivity of animal fecal DNA with Bacteroidales PCR markers, especially for the human HF183 marker. This occurred for some known source wildlife samples in the Buck Creek project (TSSWCB project 06-11) which were collected from a remote site which had very limited human access. This may explain the unexpected and frequent occurrence of water samples positive for the human marker at this site. To help explore the issue of cross-reactivity, all 100 known source fecal samples collected under this project will be analyzed for the human HF183 marker. Further development and evaluation of these library-independent methods will be conducted for possible inclusion into Texas' BST toolbox.

Due to the current and anticipated need for BST studies in Texas, statewide BST analytical infrastructure needs to be maintained appropriately. This not only includes the needed maintenance and repairs of analytical equipment; but also the continued support, training, and retention of skilled personnel. To meet the needs of the State, BST analytical capabilities will be maintained at both UTSPH EP and AgriLife SCSC BST laboratories. Financial support will be used to hire and train graduate students or a postdoctoral student at UTSPH EP and retain (or hire) graduate students or a postdoctoral student at UTSPH EP and retain (or hire) graduate students or a postdoctoral associate at AgriLife SCSC. Training needs for each individual laboratory's personnel will be coordinated to ensure appropriate technology transfer and comparability of BST data.

Delivering educational and informational programming regarding BST is also a critical need. Although the Task Force recommended the usage of BST, the TSSWCB and TCEQ adopted the general process laid out by the Task Force on the use of BST, and BST has been successfully employed in many watersheds across the state, BST is still not being used to its full potential in Texas. To provide greater outreach to water resource managers in Texas, the project team

will participate in conferences including the 2013 and 2014 TCEQ Environmental Trade Fair and Conference and other events in Texas. Flyers, one-pagers, tri-folds or other appropriate printed media developed through previous projects will be used to 1) describe the general use of BST consistent with the Task Force Report, 2) discuss the appropriate application of BST in identifying fecal contamination sources, and 3) review the analytical lab capability of public BST labs which the state has invested.

TWRI will continue to host and maintain the BST website (<u>http://texasbst.tamu.edu/</u>) to disseminate educational materials, project updates, science updates, and other outreach efforts to advance the science and application of BST in Texas and nationally.

This project will advance the recommendations of the Task Force by updating, expanding, and prioritizing BST-related R&D needs. Additionally, this project will work towards accomplishing R&D needs identified in the Report:

- Investigation and refinement of library-independent BST methods, and determine which library-independent BST methods are best suited for Texas. Specifically, this will include work on feral hog, poultry, and human markers.
- Continue expansion and refinement of the Texas E. coli BST Library.
- Continued investigations into the geographic stability of the Texas *E. coli* BST Library and refinement of library isolate selection.

While previous studies have utilized appropriate quality assurance and quality control mechanisms as identified in project-specific QAPPs, the volume of current and anticipated BST studies across the State favors the development and implementation of BST template-SOPs. BST template-SOPs developed under TSSWCB projects 08-50 and 08-51 have provided for the continued development and use of the Texas *E. coli* BST Library by multiple laboratories and will also support and improve inter-laboratory comparison of BST results. In this project, ERIC-PCR, RP and *Bacteroidales* PCR template-SOPs will be reviewed and updated accordingly to ensure that they are current and up to date with applicable methods, technologies and markers.

In order to reduce bacteria and other pollutant contributions to streams, TWRI will also coordinate a Southwestern United States Stream Conference Workshop titled: Riparian Vegetation Workshop – Putting the 'green' into streambank stabilization in San Antonio in 2013. Establishment of riparian vegetation is one of the most important components of streambank stabilization and stream restoration, but it can also be one of the most challenging. This informative half day workshop will focus on the role of riparian vegetation, overcoming challenges of riparian management and restoration, and methods of establishment. It will also discuss budgeting for and monitoring of riparian vegetation restoration efforts and the techniques for managing invasive species. Instructors for the workshop will be from multiple states including Arkansas, Oklahoma, New Mexico and Texas. Further, with assistance from the USDA-NASS Texas Field Office, a stratified random sampling scheme will be implemented to support assessment of barriers to bacteria BMP adoption in conjunction with TSSWCB Project #12-08.

Project Goals (Expand from Summary Page)

Support BST analyses across the State through (1) continued personnel support and operation and maintenance of analytical infrastructure at public BST laboratories; (2) continued development, updating and implementation of statewide BST template-SOPs for ERIC-PCR, RiboPrinting, and *Bacteroidales* PCR along with coordination amongst other entities conducting BST in the state to standardize methodologies employed; (3) delivery of information on BMPs and materials that give an overview of BST activities in Texas to date and describe the use, capabilities and applicability of BST and the services provided by the State-supported analytical labs to local, state and national stakeholder audiences; (4) continued development of the Texas *E. coli* BST Library by incorporating additional known source fecal sample isolates; and, (5) further development of suitable source-specific bacteria markers for library independent BST.

Measures of Success (Expand from Summary Page)

- Updated BST template-SOPs for ERIC-PCR, RiboPrinting, and *Bacteroidales* PCR ensuring that template-SOPs include current methods, technologies and approaches.
- Maintain needed level of training of AgriLife SCSC and UTSPH EP personnel.
- Continued operation and maintenance of BST analytical equipment and support of personnel needs to sustain operating capability and expand the utilization of BST applications statewide.
- Data analysis for approximately 100 known source *E. coli* isolates from the Leona River (TSSWCB project 11-50) for expansion of the Texas *E. coli* BST Library
- Fingerprinting and analysis of 20 known-source *E. coli* isolates collected as part of TSSWCB project 11-51 *Instream Bacteria Influences from Bird and Bat Habitation of Bridges*
- Expansion of the Texas *E. coli* BST Library through the analysis of approximately 100 known source fecal samples collected by IRNR
- Evaluation of geographical and temporal stability of the Texas *E. coli* BST Library and diversity of source specific isolates
- Development/evaluation of new source-specific bacterial markers (e.g., poultry, feral hog from domestic swine,deer from other ruminants) for library-independent BST
- Continued outreach through a BST state of the science website (<u>http://texasbst.tamu.edu/</u>) that serves as a repository for collected/produced BST information and source of BST related materials, updates, meeting announcements for educational opportunities
- Continued outreach through delivery of BST and BMP informational materials describing the state of the science, applicability, usefulness, and analytical capabilities of State-supported BST laboratories to water resource professionals across the state and nation

2012 Texas NPS Management Program Reference (Expand from Summary Page)

Components, Goals, and Objectives

Component 1 - Explicit short- and long-term goals, objectives, and strategies that protect surface... water.

LTG 1 – Objective 1 – Focus ... available resources in watersheds and aquifers identified as impacted by NPS pollution LTG 1 – Objective 2 – Support the implementation of state, regional, and local programs to prevent NPS pollution through assessment...

LTG 1 – Objective 3 – Support the implementation of state, regional, and local programs to reduce NPS pollution, such as the implementation of strategies defined in TMDL I-Plans, [and] WPPs...

LTG 1 – Objective 6 – Develop partnerships ... to facilitate collective, cooperative approaches to manage NPS pollution.

Short-Term Goal One – Data Collection and Assessment – Objective C – Conduct special studies to determine sources of NPS pollution and gain information to target... BMP implementation.

Component 2 – Working partnerships and linkages to appropriate State, interstate, Tribal, regional, and local entities, private sector groups, and Federal agencies.

Component 3 – Balanced approach that emphasizes both statewide NPS programs and on-the-ground management of individual watersheds.

Component 5 – ... Progressively address these identified waters by conducting more detailed watershed assessments...

References

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- Bernhard, A. E. and K. G. Field (2000). "A PCR assay to discriminate human and ruminant feces on the basis of host differences in Bacteroides-Prevotella genes encoding 16S rRNA." <u>Appl Environ Microbiol</u> **66**(10): 4571-4574.
- Casarez, E. A., S. D. Pillai, et al. (2007). "Direct comparison of four bacterial source tracking methods and a novel use of composite data sets." J Appl Microbiol **103**(2): 350–364.
- Dick, L. K., A. E. Bernhard, et al. (2005). "Host distributions of uncultivated fecal Bacteroidales bacteria reveal genetic markers for fecal source identification." <u>Appl Environ Microbiol</u> **71**(6): 3184-3191.
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- Field, K. G., E. C. Chern, et al. (2003). "A comparative study of culture-independent, library-independent genotypic methods of fecal source tracking." J Water Health 1(4): 181-94.
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- USEPA (2005). <u>Method 1603</u>: <u>Escherichia coli (E. coli) in water by membrane filtration using modified membrane-thermotolerant</u> Escherichia coli agar (Modified mTEC). Washington, DC, Office of Research and Development, Government Printing Office.

Weidhaas, J. L., T. W. Macbeth, et al. (2010). "Identification of a *Brevibacterium* marker gene specific to poultry litter and development of a quantitative PCR assay." J. Appl. Microbiol. 109:334-347.

| Tasks, Objec | tives and Schedules | | | | | |
|--------------|--|------------------------------|------------------------------|-----------------------------------|--|--|
| Task 1 | Project Administration | | | | | |
| Costs | \$20,000 | | | | | |
| Objective | To effectively administer, | coordinate and monitor al | l work performed under thi | s project including | | |
| | technical and financial sup | pervision and preparation of | of status reports. | | | |
| Subtask 1.1 | TWRI will prepare electro | onic quarterly progress repo | orts (QPRs) for submission | to the TSSWCB. QPRs | | |
| | shall document all activiti | es performed within a quar | ter and shall be submitted | by the 15 th of March, | | |
| | June, September, and Dec | ember. QPRs shall be distr | ibuted to all Project Partne | ers and posted on the | | |
| | project website. | | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | | |
| Subtask 1.2 | TWRI will perform accou | nting functions for project | funds and will submit appr | ropriate Reimbursement | | |
| | Forms to TSSWCB at lease | st quarterly. | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | | |
| Subtask 1.3 | TWRI will host coordinat | ion meetings or conference | e calls with the TSSWCB, 1 | UTSPH EP, and AgriLife | | |
| | SCSC at least quarterly to | discuss project activities, | project schedule, communi | cation needs, | | |
| | deliverables, and other rec | uirements. TWRI will dev | elop lists of action items n | eeded following each | | |
| | project coordination meet | ing and distribute to projec | t personnel. | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | | |
| Subtask 1.4 | TWRI will work with Ag | iLife SCSC and UTSPH E | P to develop a Final Repor | t that summarizes | | |
| | activities completed, conc | lusions reached during the | project, and the extent to v | which project goals and | | |
| | measures of success have | been achieved. | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | | |
| Deliverables | • QPRs in electronic for | nat | | | | |
| | Reimbursement Forms | , and necessary supporting | documentation, in hard coj | py format | | |
| | • Final Report in electronic and hard conv formats | | | | | |

| Tasks, Objec | tives and Schedules | | | | | | |
|--------------|--|-----------------------------|--|--|--|--|--|
| Task 2 | Quality Assurance | | | | | | |
| Costs | \$5,000 | | | | | | |
| Objective | Develop and implement d | ata quality objectives (DQ | Os) and quality assurance/o | control (QA/QC) | | | |
| | activities to ensure data of | f known and acceptable qua | ality are generated through | this project. Update and | | | |
| | implement statewide BST | template-SOPs. | | | | | |
| Subtask 2.1 | TWRI will work with UT | SPH EP, AgriLife SCSC, a | and IRNR to develop a QA | PP for activities in Tasks | | | |
| | 3-4 consistent with EPA F | Requirements for Quality A | ssurance Project Plans (Q | <i>A</i> / <i>R</i> -5) (May 2006) and | | | |
| | the TSSWCB Environmen | tal Data Quality Managem | ent Plan (August 2007). | | | | |
| | Start Date | Month 1 | Completion Date | Month 3 | | | |
| Subtask 2.2 | TWRI will submit revisio | ns and necessary amendme | ents to the QAPP as needed | l | | | |
| | Start Date | Month 4 | Completion Date | Month 24 | | | |
| Subtask 2.3 | AgriLife SCSC and UTSI | PH EP will maintain and up | odate, at least annually, the | 7 statewide BST | | | |
| | template-SOPs for collect | ion of fecal samples for BS | ST, isolation of <i>E. coli</i> , arcl | hival of E. coli isolates, | | | |
| | ERIC-PCR, RP, pre-proce | essing of water samples for | Bacteroidales PCR, and B | <i>Bacteroidales</i> PCR | | | |
| | consistent with EPA Guid | lance for Preparing Standa | ord Operating Procedures (| (SOPs) (QA/G-6) and the | | | |
| | TSSWCB Environmental | Data Quality Management | <i>Plan</i> so that they include the | he most recent advances | | | |
| | in BST science, methodol | ogies, markers and technol | ogies. | 1 | | | |
| | Start DateMonth 1Completion DateMonth 24 | | | | | | |
| Subtask 2.4 | AgriLife SCSC and UTSPH EP will coordinate to ensure that needed personnel training is kept on par | | | | | | |
| | between the groups to ens | ure congruity statewide. | | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | | | |

| Subtask 2.5 | UTSPH EP and AgriLife SCSC will work with public and private laboratories across the state which are | | | | | | | |
|--------------|--|------------------|-----------------|----------|--|--|--|--|
| | exploring the use of BST. UTSPH EP and AgriLife SCSC will work to ensure that methodologies and | | | | | | | |
| | QA/QC mechanisms adopted by these other laboratories are as congruent as possible with SOPs utilized | | | | | | | |
| | by UTSPH EP and AgriLife SCSC (subtask 2.1). | | | | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | | | | |
| Deliverables | QAPP for Tasks 3-4 approved by TSSWCB in both electronic & hard copy formats | | | | | | | |
| | Approved revisions and amendments to QAPP | | | | | | | |
| | • Updated statewide BS | ST template-SOPs | | | | | | |

| Tasks, Objec | tives and Schedules | | | | | | |
|--------------|---|-----------------------------|------------------------------|---------------------------|--|--|--|
| Task 3 | Known Source Fecal Sample Collection | | | | | | |
| Costs | \$30,000 | • | | | | | |
| Objective | To expand the Texas <i>E. co</i> fecal samples. | oli BST Library through th | e collection of approximate | ely 100 known source | | | |
| Subtask 3.1 | TWRI will work with IRNR to collect known source fecal samples. | | | | | | |
| | Start Date | Month 1 | Completion Date | Month 2 | | | |
| Subtask 3.2 | TWRI and IRNR will w | ork with UTSPH EP and | d AgriLife SCSC to deve | elop a targeted list of | | | |
| | needed species/watersh | eds for fecal sample colle | ection and plan for their | collection and delivery. | | | |
| | This list should primari | ly fill gaps in the Texas I | E. coli BST Library ident | tified in other | | | |
| | TSSWCB-funded BST | projects. Targeted specie | es will include small man | nmals such as mice, | | | |
| | squirrels, nutria and rab | bits. In addition, samples | s will be collected from a | at least one previously | | | |
| | studied watershed (e.g., | Leon River) in order to | determine the temporal s | stability of the Texas E. | | | |
| | coli BST Library. Appr | oximately 50 known sou | rce fecal samples from e | ach of 2 watersheds | | | |
| | (Leon and San Antonio | Rivers) are budgeted for | r collection (total of 100 | samples). TWRI, | | | |
| | UTSPH EP, and AgriLi | fe SCSC will review the | draft QAPP with IRNR | and discuss and | | | |
| | resolve issues as necess | ary. | | | | | |
| | Start Date | Month 2 | Completion Date | Month 4 | | | |
| Subtask 3.3 | IRNR will collect fecal | samples in accordance w | with the plan developed in | n Subtask 3.2 and work | | | |
| | closely with UTSPH EI | P and AgriLife SCSC to | coordinate delivery of the | e samples to the | | | |
| | appropriate lab. IRNR v | will communicate with a | select group of organization | tions, agencies and | | | |
| | businesses in each of th | e 2 targeted watersheds t | to arrange and resolve an | y access concerns and | | | |
| | gather input to improve | geographic targeting of | sample collection. Trave | l plans, scheduling, | | | |
| | and routing maps will b | e prepared prior to deplo | ying the field crew. IRN | R will deploy the field | | | |
| | crew to collect known s | source samples from each | n targeted watershed. IRN | NR will coordinate | | | |
| | closely with UTSPH EI | P and AgriLife SCSC to | ensure sample delivery a | dheres to established | | | |
| | QA/QC procedures. A l | known source sample dat | ta set will be finalized af | ter completion of the | | | |
| | field work and submitte | d to TWRI. | | | | | |
| | Start Date | Month 4 | Completion Date | Month 15 | | | |
| Deliverables | • Map of watersheds targ | geted for known source san | nple collection | | | | |
| | Proposed list of needed | l species recommended for | fecal sample collection | | | | |
| | • MS Excel summary da | ta sheets cataloguing know | n source samples collected | l | | | |

| Tasks, Objec | tives and Schedules | | | | |
|--------------|--|--------------------------------|-------------------------------------|-----------------------------|--|
| Task 4 | Analytical Laboratory Ca | pacity Library Expansion | and Methods Developmen | t | |
| Costs | \$205.348 | | | | |
| Objective | Support BST analyses across Texas, through continued operation and maintenance of BST laboratory | | | | |
| | analytical infrastructure including equipment and personnel. Evaluate and expand the statewide <i>E</i> coli | | | | |
| | BST library through the analysis of ERIC-RP data provided by AgriLife SCSC for approximately 100 | | | | |
| | <i>E. coli</i> known source isola | ates obtained from the Leon | na River watershed (TSSW | CB Project 11-50) and | |
| | the addition of known sou | rce fecal samples collected | through Task 3 and TSSV | VCB project 11-51. | |
| | Develop and refine library | -independent markers. | | r J | |
| Subtask 4.1 | UTSPH EP and AgriLife | SCSC will maintain BST a | nalytical equipment (e.g., l | RiboPrinter) and general | |
| | laboratory equipment. Thi | is includes securing mainte | enance contracts, replaceme | ent parts, and expendable | |
| | supplies and purchase of a | a new computer for the UT | SPH EP RiboPrinter syster | n. | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtask 4.2 | UTSPH EP will retain (or | hire) a Graduate Student c | or Postdoctoral Research As | ssociate that will 1) | |
| | maintain laboratory opera | ting capacities and technica | al expertise to conduct BST | Γ studies across the state, | |
| | 2) aid in the evaluation, ex | xpansion and maintenance | of the Texas E. coli BST L | ibrary, 3) evaluate | |
| | library-independent metho | ods and markers, and 4) pro | ovide support on TSSWCB | project 12-10 BST to | |
| | Support Adaptive Manage | ement of the Arroyo Colora | udo WPP. | | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtask 4.3 | AgriLife SCSC will retain | (or hire) Graduate Studen | ts and/or a Postdoctoral Re | esearch Associate that | |
| | will 1) maintain laboratory | y operating capacities and | technical expertise to cond | uct BST studies across | |
| | the state, 2) continue BST | efforts in support of TSSV | WCB projects 09-10 Develo | opment of a WPP for | |
| | Attoyac Bayou, 11-50 Ass | essment of Water Quality c | and Watershed Planning fo | r the Leona River, and | |
| | 11-51 Instream Bacteria I | Influences from Bird and B | <i>at Habitation of Bridges</i> , a | nd 3) evaluate new | |
| | poultry marker(s) for library-independent BST. | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtask 4.4 | UTSPH EP and AgriLife | SCSC will expand the state | ewide <i>E. coli</i> BST library th | hrough the analysis of | |
| | ERIC-RP data provided b | y AgriLite SCSC for appro | Example 11 50 E. coli know | n source isolates | |
| | obtained from the Leona Kiver watershed (155 WCB Project 11-50). Additionally, U18PH EP and | | | | |
| | AgriLite SUSU will isolate <i>E. coll</i> from approximately 100 known source fecal samples collected through Task 3, which should primarily fill gons in the library identified in other TSSWCD funded DST | | | | |
| | noisets. Approximately three isolates from each feeal sample (for a total of approx 300 isolates) will be | | | | |
| | analyzed using ERIC-PCR for inclusion in the Tayas <i>E</i> coli RST L ibrary based on the EDIC DCP | | | | |
| | fingerprint patterns approximately half of the isolates (150) will be further analyzed using RP for | | | | |
| | inclusion in the Texas <i>E. coli</i> BST Library UTSPH FP and Aoril ife SCSC will equitably solit | | | | |
| | workload. AgriLife SCSC will also fingerprint (ERIC-RP) and analyze 20 known-source <i>E. coli</i> isolates | | | | |
| | collected as part of TSSWCB Project 11-51. | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtask 4.5 | UTSPH EP and AgriLife | SCSC will collaborate to e | valuate the geographical ar | nd temporal stability, | |
| | composition, average rate | s of correct classification (a | accuracy), diversity of sour | ce specific isolates, and | |
| | further development and r | efinement needs of the Tex | xas <i>E. coli</i> BST library. | 1 | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtask 4.6 | Using known source fecal | material, AgriLife SCSC | and UTSPH EP will utilize | the best available | |
| | bacterial indicators to eval | luate and further develop/r | efine source-specific bacter | rial PCR markers. | |
| | Specifically, efforts will b | e made on markers to 1) ic | lentify poultry litter/manur | e pollution, 2) evaluate | |
| | the use of genetic targets l | based on ERIC-PCR produ | cts to differentiate human a | and animal derived | |
| | E. coli, 3) differentiate be | tween domestic swine and | feral hogs, 4) differentiate | deer from other | |
| | ruminants by continued an | nalysis of existing data on o | deer fecal microbial comm | unities, and 5) evaluate | |
| | the occurrence of human I | HF183 marker cross reactive | vity for all 100 known sour | ce animal samples | |
| | collected under Task 3. | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | |

| Subtask 4.7 | AgriLife SCSC and UTSPH EP will cooperate with other entities nationwide to ensure that the most up- | | | | | |
|--------------|--|--|--|--|--|--|
| | to-date and accurate BST approaches are implemented in Texas by attending and participating in BST- | | | | | |
| | related meetings, seminars and workshops, as appropriate, to learn of new and improved BST methods | | | | | |
| | being employed elsewhere. | | | | | |
| | Start DateMonth 1Completion DateMonth 24 | | | | | |
| Deliverables | • Highlights of work performed included in OPRs and Final Report | | | | | |

| Tasks, Objec | tives and Schedules | | | | |
|--|--|------------------------------|------------------------------|---------------------------|--|
| Task 5 | Outreach on Bacterial Source Tracking and BMPs | | | | |
| Costs | \$65,000 | <u> </u> | | | |
| Objective | To further outreach regarding bacteria BMPs as well as the science of BST and its application through | | | | |
| 5 | improving the statewide knowledge base regarding current BST practices BMPs, scientific advance | | | | |
| | improvements in the appl | ication of BST and incorpo | orating information from ot | her areas of the nation | |
| | into the BST approaches | utilized in Texas. | C | | |
| Subtask 5.1 | TWRI will host and maintain the http://texasbst.tamu.edu website to disseminate educational materials. | | | | |
| | project updates, science u | pdates, notify readers abou | it educational opportunities | , and other outreach | |
| | efforts to advance the scie | ence and application of BS | Γ in Texas and nationally. | | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtask 5.2 | TWRI, UTSPH EP, and A | AgriLife SCSC will periodi | cally meet with natural reso | ource agencies, including | |
| | but not limited to USEPA | -R6, TCEQ, TPWD, TDA | , GLO, DSHS, and selected | l river authorities, to | |
| | advance the general know | ledge and understanding o | f agency staff on BST and | to develop action | |
| | strategies to address issue | s raised by agency staff reg | garding the use of BST in T | exas. | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtask 5.3 | Subtask 5.3 TWRI, UTSPH EP, and AgriLife SCSC will distribute the educational brochures developed t | | | | |
| | TSSWCB Project 10-50 (subtask 4.2). TWRI, UTSPH EP, and AgriLife SCSC will develop additional | | | | |
| | flyers, one-pagers, tri-folds or other appropriate printed media, as needed, that can be used to 1 | | | | |
| | the appropriate application of BST in identifying fecal contamination sources, and 2) promote the analytical laboratory capability of public BST labs which the State has invested. As appropriate, TWRI | | | | |
| | | | | | |
| | will include information about BST in general, and this project specifically, in the txH2O magazine, the | | | | |
| Conservation Matters e-mail newsletter, and AgriLite I oday news. | | | | | |
| ~ 1 1 7 4 | Start Date | Month I | Completion Date | Month 24 | |
| Subtask 5.4 TWRI, UTSPH EP, and AgriLife SCSC will promote the use of and provide resources on F | | | | sources on BST by | |
| | participating in conferences, worksnops, seminars, and other appropriate venues, including but not limited to the 2012 and 2014 TCEO Environmental Trade East, WEE/WEAT events in Taylor | | | | |
| | Imited to the 2013 and 2014 TCEQ Environmental Trade Fair, wEF/wEAT events in Texas, | | | | |
| | ISCRA/IFB/IWA annua | al conventions, and ASABI | E events in Texas. | Mandh 24 | |
| Subtack 5.5 | TWDI LITSDILED and A | MONIN I | inform other records and | Month 24 | |
| Sublask 5.5 | .5 I WKI, UI SPH EP and AgriLite SUSC will work to inform other researchers/academia who are | | | cademia who are | |
| | engaged in BS1 in Texas (e.g., Edrington, Brinkmeyer, Alam, Ward) about the methods and approach | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| Subtack 5.6 | To build on the success of | f the 2012 RST State of the | ha Science Conference TW | /PI LITSPH FP and | |
| Sublask 5.0 | A gril if a SCSC will evolute the mood for and timing of a fallow we conference. I WKI, UISPH | | | If the need is | |
| | substantiated TWRI LITSPH EP and Agril if SCSC will initiate planning and logistics for a fallow | | | | |
| | conference | | | | |
| | Start Date | Month 1 | Completion Date | Month 24 | |
| | Start Date | | Completion Date | Monul 24 | |

| Subtask 5.7 | With assistance from the USDA-NASS Texas Field Office, a stratified random sampling scheme will be | | | | | | |
|--------------|---|------------------------------|--------------------------------|----------------------------|--|--|--|
| | implemented using a target population of beef cattle producers who completed 2012 Census of | | | | | | |
| | Agriculture forms. The sa | mple will be stratified acco | ording to NASS district and | l beef cattle herd size. | | | |
| | USDA-NASS will provide Texas A&M Department of Soil & Crop Sciences with a list of unique | | | | | | |
| | identifying numbers that v | will be placed on all survey | materials so that response | non-response can be | | | |
| | tracked. The USDA-NAS | S Texas Field Office will a | lso assist with logistics rela | ated to compiling, | | | |
| | stuffing, and mailing surv | ey materials that will inclu | de an introductory postcare | l, the first survey packet | | | |
| | with cover letter and survey instrument, a reminder postcard, and a second survey packet with cover | | | | | | |
| | letter and survey instrument. This information will support assessment of barriers to BMP adoption in | | | | | | |
| | conjunction with TSSWCB Project #12-08. | | | | | | |
| | Start DateMonth 1Completion DateMonth 24 | | | | | | |
| Subtask 5.8 | In order to reduce pollutant contributions to streams, including bacteria, TWRI will coordinate a | | | | | | |
| | Southwestern United States Stream Restoration Conference Workshop titled: Riparian Vegetation | | | | | | |
| | Workshop – Putting the 'green' into streambank stabilization in San Antonio in 2013. | | | | | | |
| | Start DateMonth 1Completion DateMonth 12 | | | | | | |
| Deliverables | Summaries of outreach efforts included in QPRs and Final Report | | | | | | |

Part III – Financial Information

| Budget Summary | | |
|--------------------------------|---------------|--|
| Category | Costs | |
| Personnel | \$ 125,014 | |
| Fringe Benefits | \$ 34,735 | |
| Travel | \$ 15,237 | |
| Equipment | \$ 0 | |
| Supplies | \$ 12,385 | |
| Contractual | \$ 160,600 | |
| Construction | \$ 0 | |
| Other | \$ 29,945 | |
| | | |
| Total Direct Costs | \$ 377,916 | |
| Indirect Costs ($\leq 15\%$) | \$ 37,432 | |
| | | |
| Total Project Costs | \$ 415,348 | |

| Budget Justification | | |
|----------------------|-------------------|--|
| Category | Total Amount | Justification |
| Personnel | \$ 125,014 | • TWRI Associate Director (0.1 FTE) = \$16,485 |
| | | • SCSC Assistant Professor (0.08 FTE) = \$17,499 |
| | | • IRNR Website Administrator $(0.04 \text{ FTE}) = \$3,341$ |
| | | • IRNR Extension Assistant (0.16 FTE) = \$6,839 |
| | | • IRNR Research Scientist (0.16 FTE) = \$9,558 |
| | | • Postdoctoral Associate (0.8 FTE) = \$64,960 |
| | | • Undergraduate Student Labor (0.15 FTE) = \$6,332 |
| Fringe Benefits | \$ 34,735 | Calculated at 17.2% of Personnel (9.9% for Graduate Students) to cover |
| | | FICA, UCI, WCI, and retirement. Additional \$474/mo. (\$376/mo. For |
| | | Graduate Students) prorated per %FTE is calculated for group health |
| TT 1 | ф. 15.007 | Insurance. |
| Travel | \$ 15,237 | • TWRI Associate Director (\$1,144) |
| | | $\circ \text{Per diem ($284)}$ |
| | | • Lodging ($\frac{1}{216}$) • Miles es (1.160 mil @ \$0.555/mil = \$(44) |
| | | O Mileage (1,100 mil. (a) \$0.555/mil = \$644) |
| | | • IRINK Extension Assistant & Research Scientist (54,000) |
| | | • Instructor Travel – Southwest Stream Restoration Conference (\$4,845) |
| | | • SUSC Assist. Prof. & Grad Students (\$5,250) |
| | | National Meetings (\$5,750) State Meetings (\$1,500) |
| Fauinment | \$ 0 | N/A |
| Supplies | \$ 12 385 | $\mathbf{I}_{\mathbf{N},\mathbf{A}}$ |
| Supplies | φ 12,505 | • SCSC supplies = $$11,885$ |
| | | Computer for postdoc = \$1,000 |
| | | \sim FRIC-RP Supplies for project 11-51 (\$53 x 20 isolates) = \$1.060 |
| | | \odot E coli isolation/archival from fecal samples (\$25 x 50) = \$1,000 |
| | | • ERIC-RP Supplies for new projects (\$8 x 150 ERIC: \$45 x 75 RP) |
| | | = \$4,575 |
| | | Eval/Development Supplies = \$4,000 |
| Contractual | \$ 160,600 | • UTSPH $EP = $153,371$ |
| | | • USDA-NASS = \$7,229 |
| Construction | <u>\$</u> 0 | N/A |
| Other | \$ 29,945 | • Booths at Environmental Trade Fair = \$1,700 |
| | | RiboPrinter Preventative Maintenance and Service (for RiboPrinters at |
| | | both UTSPH EP and AgriLife SCSC) (TWRI) = \$15,000 |
| | | • Instructor fees for Southwest Stream Restoration Conference = $$2,9/5$ |
| | | • DNA Sequencing for library independent markers (SCSC) = $$2,000$ |
| | | • Conference Registration (SCSC) = $\$1,050$ |
| | | • General Maintenance on equipment $(SCSC) = $1,500$ |
| | | • NELAP Lab accreditation fees (SCSC) = $$2,020$ |
| | | • IRNR Shipping = $$500$ |
| | | • BST Brochures = $\$1,000$ |
| T 1' / | φ 2= 125 | • Journal Publication Charges = \$2,200 |
| Indirect | \$ 37,432 | 15% of Modified Total Direct Costs (Total minus Contractual >\$25,000 per |
| SOURCE | TSSWCD will me | i contract and Equipment) vide \$415,348 in non-federal funds sourced from state appropriations |
| SOURCE | (EV2013 Conorol) | viue \$413,546 III non-rederat runds sourced from state appropriations |
| | (1°12013 General | Kevenue) unough the wonpoint Source Orant Program to 1 WKI. |

| Contractual Budget Justification – UTSPH EP | | | |
|---|-------|--------|---|
| Category | Total | Amount | Justification |
| Personnel | \$ | 66,736 | • El Paso: Di Giovanni, PI (0.1 FTE) = \$26,136 |
| | | | • El Paso: Grad Student (0.50 FTE) = 40,600 |
| Fringe Benefits | \$ | 15,205 | • El Paso: Di Giovanni @ 24% of personnel = \$6,273 |
| | | | • El Paso: Grad Student @ 22% of personnel = \$8,932 |
| Travel | \$ | 8,000 | • Di Giovanni and Grad Student/Postdoc = \$8,000 |
| | | | • National Meetings $(\$3,000/\text{yr}) = \$6,000$ |
| | | | • State Meetings $(\$1,000/yr) = \$2,000$ |
| Equipment | \$ | 9,000 | Computer & software for DuPont Qualicon RiboPrinter |
| Supplies | \$ | 30,825 | • E. coli isolation and archival from known source fecal samples (\$25 x |
| | | | 50) = \$1,250 |
| | | | • ERIC-RP supplies (\$8 x 150 ERIC, \$45 x 75 RP) = \$4,575 |
| | | | • Bacteroidales human HF183 PCR analysis of fecal samples (\$75 x 100) = \$7,500 |
| | | | • Supplies for library independent method eval/development, sequencing = \$17,500 |
| Contractual | \$ | 0 | N/A |
| Construction | \$ | 0 | N/A |
| Other | \$ | 3,600 | • Meeting registration fees = \$1,600 |
| | | | • General Maint.(Biological Safety Cabinet, freezers and refrigerators) = |
| | | | \$2,000 |
| Indirect | \$ | 20,005 | 15% of Modified Total Direct Costs |