I. Results

The hypothesis of our research project was that microbial communities respond to sub-toxic heavy metal stress by increased horizontal gene transfer and rearrangement. We postulated that such response leads to useful genetic rearrangements and recombinations that improve the community's ability to resist or cope with the applied heavy metal stress. Our thrust was, therefore, to directly examine gene transfer in soil microcosms subject to different levels of heavy metal stress and examine this for soils that were either pristine or had historic heavy metal contamination.

In our microcosm experiments we mimicked the conditions of a semi-continuous oligotrophic carbon flux that exists in the subsurface. Samples were obtained from the saturated zone of an alluvial deposit aquifer at a site contaminated with Cd and Zn adjacent to a closed metal plating waste lagoon. The actual total and NaNO₃ exchangeable metal concentrations were 79 microg/g Cd, 96 microg/g Ni, 63microg/g Zn; and 35microg/g Cd, 8microg/g Ni, and 5 microg/g Zn respectively. In addition, our approach permitted facile periodic sampling of the microbial community. In brief, each microcosm consisted of subsurface soil aliquots placed in 40-ml sample vial subject to periodic (U/week) addition and withdrawal of artificial groundwater (AGW) supplemented with a low carbon concentration (peptone at 10 mg/L) to maintain a low carbon flux in the system. The withdrawn samples were then subject to molecular analysis and microbial enumeration. Repeated addition and withdrawal of AGW allow long-term (10 to 30+ weeks) operation of the microcosm. In most experiments, Cd was studied as model heavy metal stress at different concentrations (C,Cd = 0,10,100,1000 microM). All treatments were conducted in duplicate and contained controls to assess background contribution of the indigenous community to various analyses.

Several long-term microcosm experiments were completed. In a first set, the fate of Pseudomonas putida KT2440 and its plasmid TOL::Tn5npt was monitored at various degrees of Cd stress. This strain was originally proposed as the model delivery system for our work, because it contains a chromosomal copy of the P. aeruginosa gef-based IPTG inducible suicide gene cassette present on a miniTn5. Hence, the efficacy of suicide induction in the microcosm systems was examined. Our results indicated, however, that the P. aeruginosa – gef suicide miniTn5 construct was not an effective means of selective elimination of a bacterial plasmid donor in tested soil microcosms, due to the combined effects of spontaneous loss of the mini-transposon, spontaneous mutations in the P. aeruginosa – gef cassette and rapid selection of escape mutants upon suicide induction. Due to the inefficacy of the suicide system for selective plasmid donor removal, several Escherichia coli strains were subsequently used for plasmid delivery in the microcosm given their presumed competitive inferiority.
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Using subsurface soils historically contaminated with Zn$^{2+}$, Cd$^{2+}$, and Ni$^{2+}$, we examined the fate of two RSFI101 plasmids (IncQ, tra- mob+; pMOL 187 and pMOL 222 containing the czc and ncc operons, respectively) originally introduced with E. coli donors in semi-continuously operated microcosms at increasing Cd concentrations. Further, we examined whether the long-term stability of the plasmid in the community was enhanced by the simultaneous co-introduction of the broad-host range plasmid RP4 (IncPΔ, tra+ mob+). Hence, all treatments were examined in absence or presence of RP4. While the stability of the introduced E. coli strain (pMOL222 or pMOL187 result in low level heavy metal resistance in E. coli) was somewhat impacted by the degree of metal contamination, the strains were typically not recoverable after 4 weeks of operation. Total culturable cell counts remained in the range of 10$^5$ to 10$^7$ cells/ml throughout the microcosm operation. Increases in the degree of heavy metal resistance were observed only at the highest Cd addition suggesting little effect of the applied stress on the community phenotype below 1 mM Cd. No clear phenotypic evidence of pMOL 187 or pMOL 222 transfer could not be obtained: increases in Cd or Ni resistance were associated with applied Cd dose rather than the addition of the plasmids. Thus, ID-PCR was employed to enumerate the abundance of czc harbored on pMOL187 and ncc harbored on pMOL222. We obtained an ID-PCR detection limit of 1.2 $10^3$ and 1.6 $10^5$ copies in a background of $10^8$ cells for czc and ncc, respectively. Further, introduced E. coli strains were detected employing ID-PCR targeting rrsA3 (E. coli MG1655) that permitted specific enumeration of enteric bacteria. RP4 was detected employing IncPrep specific primers which target the replication function tra$^A_1$. Simultaneous RP4 introduction consistently enhanced pMOL187 and pMOL222 stability in the microbial community, suggesting that the plasmids were mobilized by the introduced RP4 plasmid. Transiently (2 to 3 weeks after introduction) high titers of $10^8$ and $10^9$ templates per ml were measured for pMOL187 and pMOL222, respectively. Further, czc was still observed in microcosms after 32 weeks ($10^7$ per ml) but only if RP4 was co-introduced, although ID-PCR with IncPrep specific primers for RP4 did not reveal continued persistence of RP4 and also introduced E. coli strains were confirmed absent at this point. The effect of increased Cd concentrations on czc persistence was marginal. Exogenous isolation using Ralstonia metallodurans as capturing strain was performed to reisolate the IncQ plasmid from the indigenous microbial community. The plasmid was only reisolated from the microcosms containing solely E. coli (pMOL222), for which a transfer frequency of $2x10^{-7}$ (cells/ml transconjugants/cells/ml recipients) was obtained, suggesting a low mobilization potential of the soil community. Transconjugants from the indigenous community harboring pMOL222 were identified by enumeration on increasing Ni concentration and detection of ncc by filter-hybridization.

Our results have clearly indicated that plasmid transfer in oligotrophic and metal-impacted subsurface microbial communities can occur at detectable frequencies. Rather than examine transfer to co-introduced recipient strains, we have followed transfer into the indigenous community employing molecular enumeration of relevant genes The extent of plasmid transfer may be limited by the mobilization potential of the indigenous community. This is remarkable, since our phenotypic community assessment did not reveal a clear benefit of the introduced plasmid. Within the range examined, metal stress did not seem to affect plasmid incidence in the community.

Several questions are raised by our research findings: What is the normal plasmid pool in the studied microbial communities, and how reflective are our model plasmids to study the dynamics of indigenous plasmids? What is the host range of our model plasmids, and what is the mobilization potential of such communities for indigenous plasmids? Is there a difference between long-term stressed and short-term perturbed microbial communities in terms of horizontal gene transfer? What molecular rearrangements occur in concordance with horizontal gene transfer, and do any of these results in adaptive traits endowed to the community? What is the physiological basis of the effect of various chemical stressors on bacterial activity and horizontal gene transfer? Answering these questions, in appropriate model systems and environments, will allow us to better ascertain the significance of horizontal gene transfer as an adaptive response in microbial communities subject to anthropogenic influences.
II. Products Delivered

Conference presentations made during the project duration that were, wholly or in part, funded by this research program:


III Notes

Several manuscripts are in advanced preparation and will be submitted with acknowledgement of funding under this research program. Preprints will be submitted to the program manager once available.

