

Extreme Spatial and Temporal Variability of Microbial Mat Communities From Mariana Island Arc Hydrothermal Vents

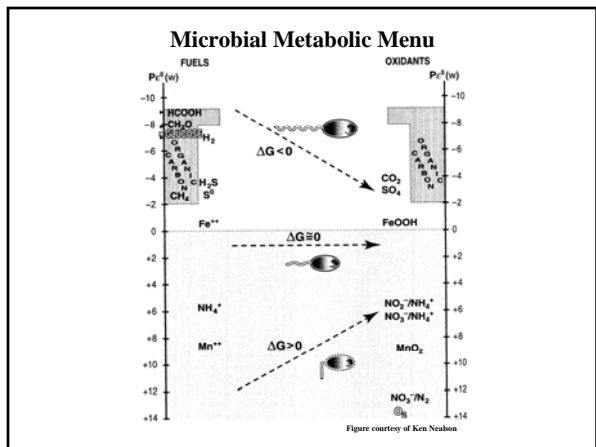
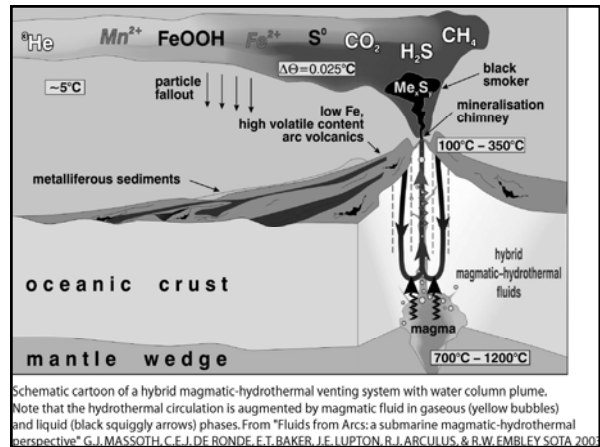
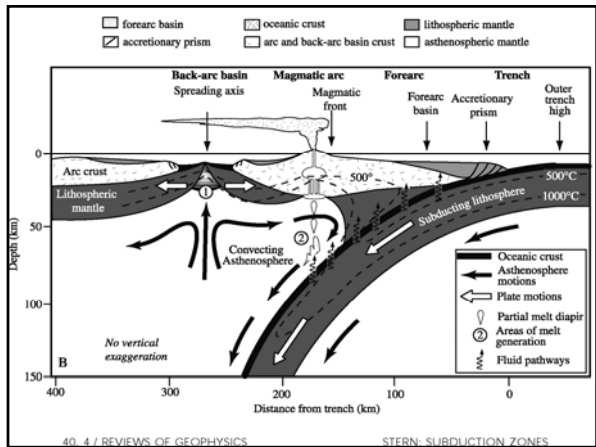
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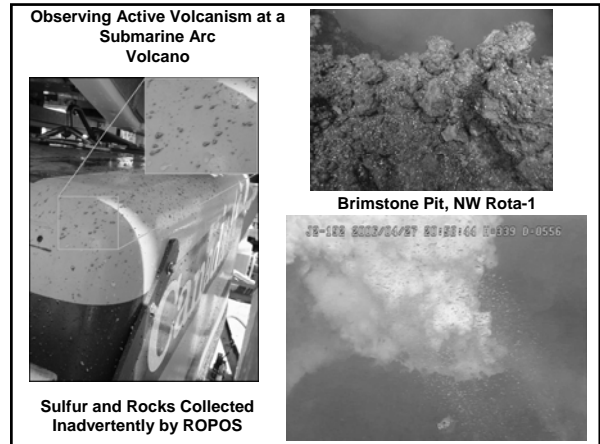
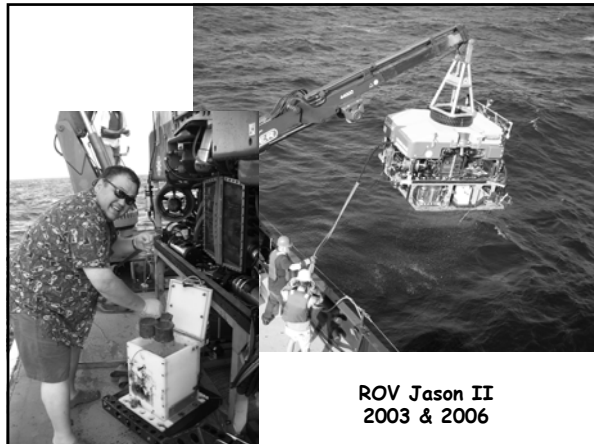
SROF Project 2003-2006

Explorer Ridge
Mariana Arc
Ring of Fire

Unlike most vent systems, the Mariana arc/backarc is a convergent plate boundary

Sample Locations





Submarine Ring of Fire Major Results/Observations

- First Observations of Deep Submarine Arc Volcanic Event
- Photosynthetic/Chemosynthetic Ecosystems Interaction
- Shallowest Massive Sulfide Formation at 344 m (220⁰ C)
- Champagne Vent, 1600 meter site with intense CO₂ Venting – Bubbles of Liquid CO₂
- High Variability of Biological Communities among & between Volcanoes

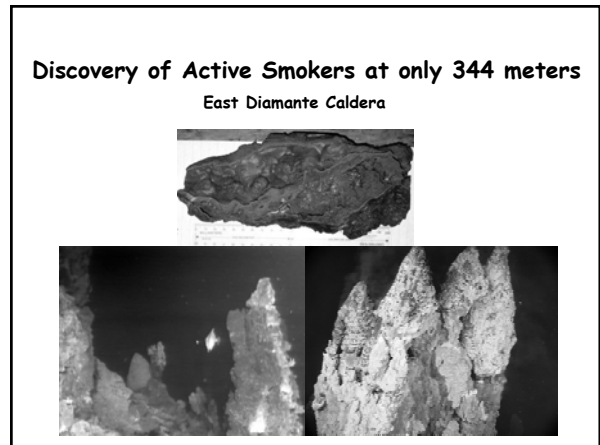


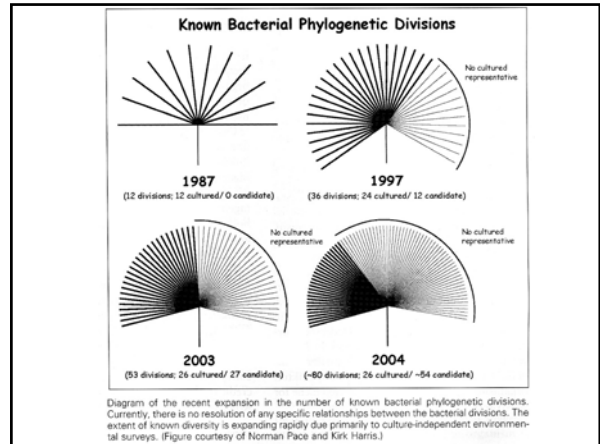
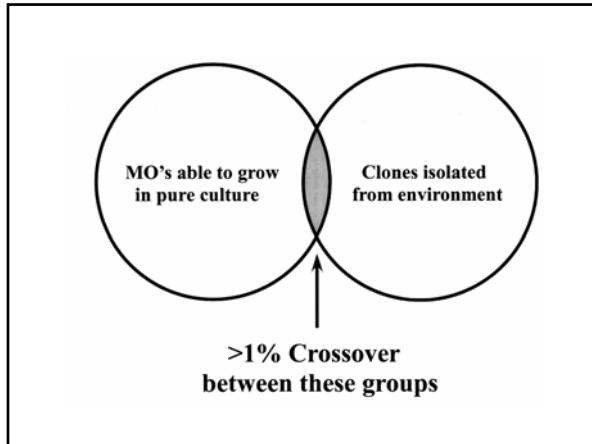
Table 1. Microbial mat samples (n=25) were collected and analyzed using T-RFLP, Q-PCR, and then targeted cloning & sequencing.

Year	Sample	Vent Site	Location	Spreading Center/ Arc Segment	Latitude/Longitude	Sample Type	Sample Description	Depth (m)	Temp (°C)
2003	JD-43-1W	Fryer Site	Backarc	SMRSC ¹	13°51.190 N 143°37.125 E	Push Core	Yellow/Green mat	2860	37
2003	JD-42-2W	Fryer Site	Backarc	SMRSC ¹	13°51.190 N 143°37.125 E	Push Core	Yellow/Green mat	2860	37
2004	R782-16	Shrimping vent	Volcanic arc	NW Rota-1	14°38.072 N 144°46.530 E	Suction Sample	Orange mat	516	15
2004	R782-17	Shrimping vent	Volcanic arc	NW Rota-1	14°38.072 N 144°46.530 E	Suction Sample	Yellow mat	518	15
2004	R783-206	Yokoh	Volcanic arc	NW Rota-1	14°38.048 N 144°46.570 E	Suction Sample	White mat	520	18
2004	R785-2537	Fault Shrimp	Volcanic arc	NW Rota-1	14°38.038 N 144°46.644 E	Suction Sample	White mat	584	20
2004	R785-2537	Fault Shrimp	Volcanic arc	NW Rota-1	14°38.032 N 144°46.518 E	Suction Sample	White mat	206	ambient
2004	R788-15	Five Towers	Volcanic arc	E Diamante	13°58.556 N 143°40.804 E	Suction Sample	Orange mat	344	220
2004	R788-22	Five Towers	Volcanic arc	E Diamante	13°58.556 N 143°40.804 E	Chimney Chunks	Multi-colored rocks	264	220
2004	R789-15	Egg Drop Ship	Volcanic arc	Mesa Crater	20°51.206 N 143°13.308 E	Suction Sample	Orange mat	344	220
2004	R789-156	Cave Vent	Volcanic arc	Mesa Crater	20°51.408 N 143°13.308 E	Suction Sample	Orange mat	145	28
2004	R791-1056	Beale Bath	Volcanic arc	NW Efluku	21°29.228 N 144°02.436 E	Suction Sample	Fe mat	1715	ambient
2004	R793-17	Yellow Top	Volcanic arc	NW Efluku	21°29.264 N 144°02.424 E	Suction Sample	Fe mat	1674	-6.8
2004	R792-1637	Champagne	Volcanic arc	NW Efluku	21°29.256 N 144°02.508 E	Suction Sample	White mat	1608	72-103
2004	R792-CC	Champagne	Volcanic arc	NW Efluku	21°29.256 N 144°02.508 E	Chimney Chunks	White rocks	1608	72-103
2004	R793-111	Yellow Cone	Volcanic arc	NW Efluku	21°29.292 N 144°02.524 E	Suction Sample	Fe mat	1587	11
2004	R793-1637	Yellow Top	Volcanic arc	NW Efluku	21°29.310 N 144°02.424 E	Suction Sample	Fe mat	1679	-6.8
2004	R793-1636	Fish Spa	Volcanic arc	Dakaku	21°18.476 N 144°11.532 E	Suction Sample	White sediment	390	ambient
2006	JD-184-W	Fe Mats	Volcanic arc	Seamount X	13°13.098 N 144°01.069 E	Suction Sample	Fe mat	1305	nd
2006	JD-184-B	Squal Mat	Volcanic arc	Seamount X	nd	Suction Sample	White mat	1188	nd
2006	JD-190-W	Fe Mounds	Volcanic arc	Esmeralda Bank	14°57.364 N 143°14.478 E	Suction Sample	Fe sediments	291	40
2006	JD-190-CC	Fe Mounds	Volcanic arc	Esmeralda Bank	14°57.364 N 143°14.478 E	Chimney Chunks	Fe crust	291	40
2006	JD-191-W	Yokoh	Volcanic arc	NW Rota-1	14°38.052 N 144°46.570 E	Suction Sample	White mat	530	25
2006	JD-191-B	Yokoh	Volcanic arc	NW Rota-1	14°38.052 N 144°46.570 E	Suction Sample	White mat	530	25
2006	JD-197-B	Rubber Bath	Volcanic arc	Dakaku	21°18.508 N 144°11.608 E	Suction Sample	White mat	411	5.2
2006	JD-197-B	Rubber Bath	Volcanic arc	Dakaku	21°18.484 N 144°11.585 E	Suction Sample	Brown sediment	390	ambient

¹ southern Mariana Backarc Spreading Center

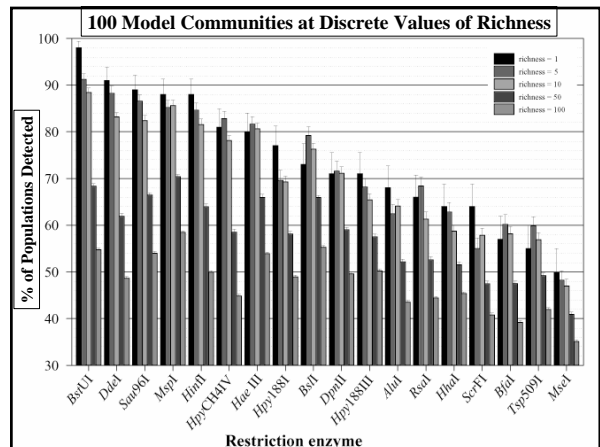
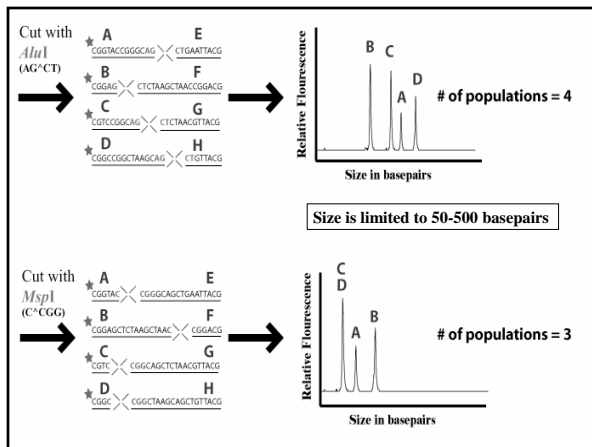
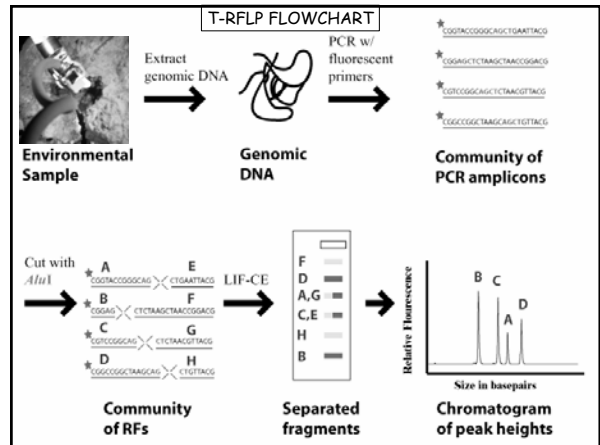
Importance of a Molecular Microbiological Approach

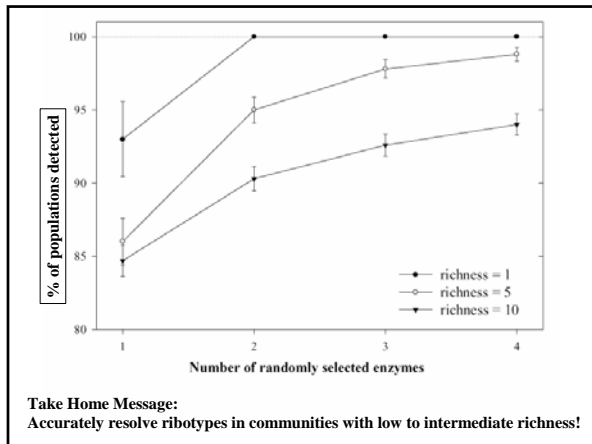
- **Traditional culturing** techniques isolate ~1% of the total bacteria in marine ecosystems, thereby severely underestimating diversity and community structure.
- Because nutrient-rich **culture media** have been historically used during enrichment procedures, bacteria which may be dominant in natural communities are selected against in favor of copiotrophic (weedy) bacteria.
- **SSU rRNAs** and their respective genes are excellent descriptors of microbial taxa based on phylogeny.



Why ribosomal RNAs?

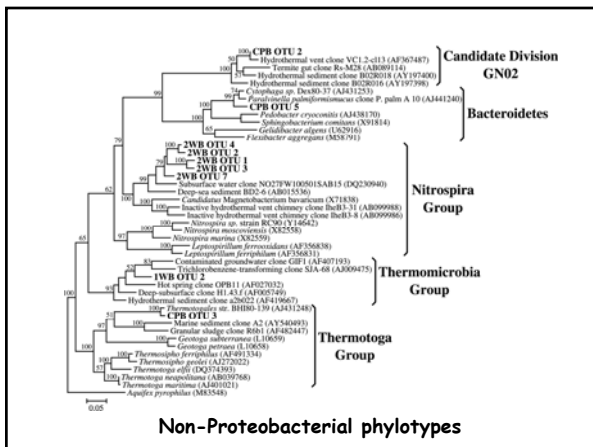
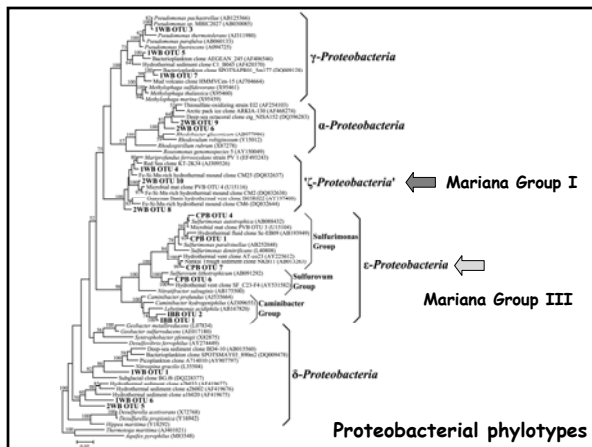
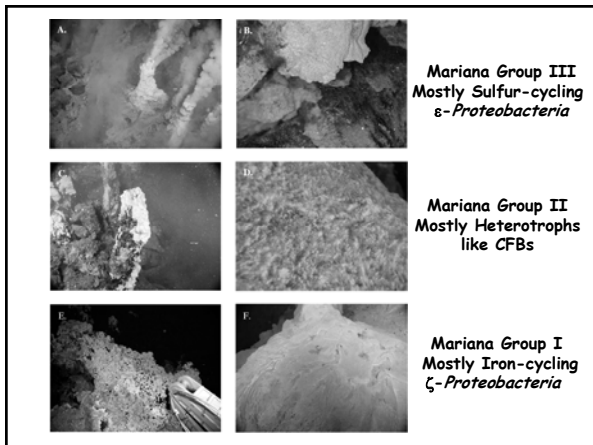
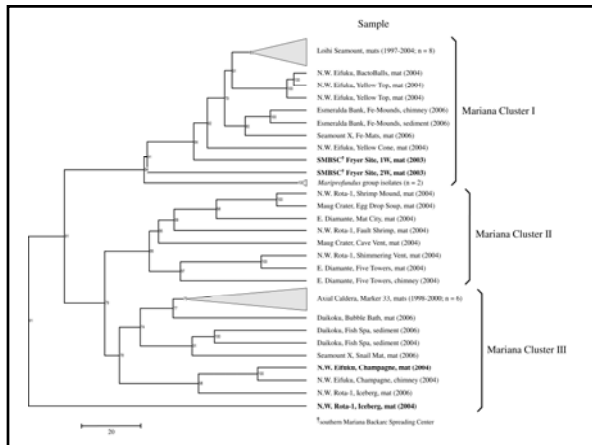
- Found among all living organisms (for 3.8 of the last 4.5 billion years). Integral part of protein synthesis machinery.
- Cell component analyses provide culture-independent means of investigating questions in microbial ecology (lack of morphology).
- rRNAs offer a type of sequence information that makes them excellent descriptors of an organism's evolutionary history.
- No detectable horizontal gene transfer, especially important for *Bacteria* and *Archaea*.
- Large and growing database; RDP contains >480K SSU rRNA genes.





Integrating Molecular Methods

- Terminal-restriction fragment length polymorphisms (T-RFLPs) were used to track populations based on ribotypes (in lots of samples).
- A/B Q-PCR needed to determine domain level proportions.
- Clone library and phylogenetic analyses needed to identify these phylotypes within the community (in few samples).

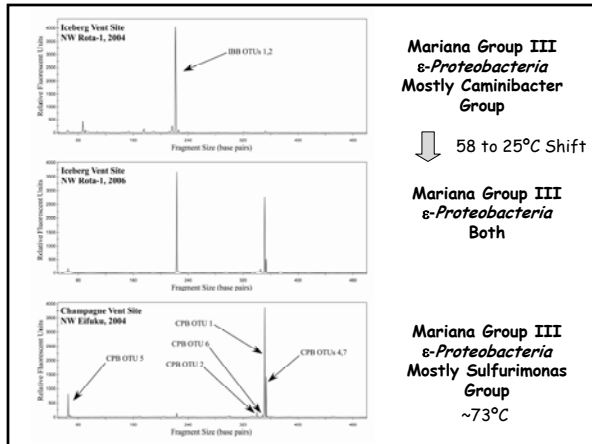
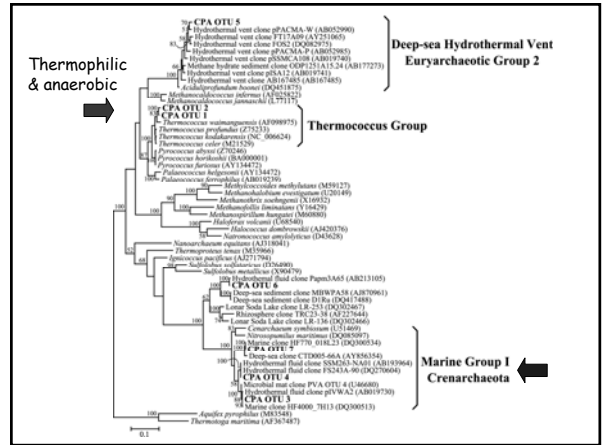
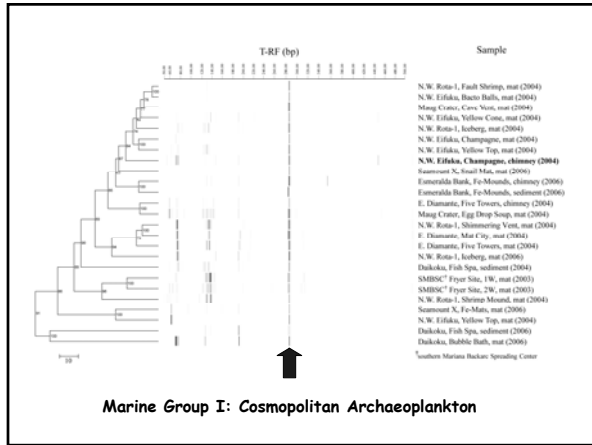
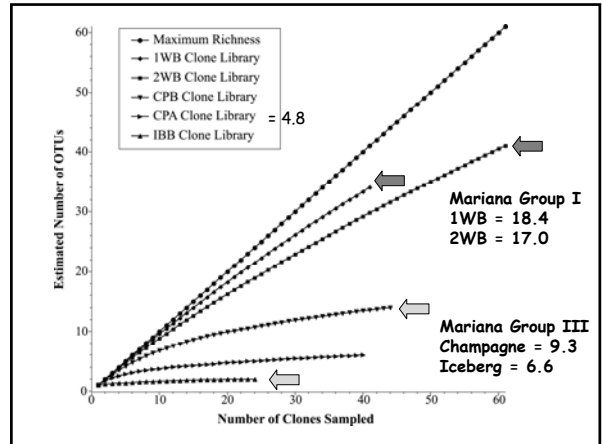


Estimate of Complexity (aka Richness)

Biomass Proxy →

Year	Sample	Vent Site	ng DNA/g Sample (wet weight)	Percent Archaea	Ave. T-RFs Bacteria	Ave. T-RFs Archaea
2003	J2-42-1W	Fryer Site	2250	11.31.49	18.444.2	18.316.5
2003	J2-42-2W	Fryer Site	2466	10.410.26	17.011.5	16.915.4
2004	R782-85	Shimmering Vent	147	2.240.16	16.643.6	9.843.5
2004	R782-87	Shrimp Mound	2450	4.440.37	12.844.0	8.313.0
2004	R783-856	Iceberg	2370	5.510.45	6.616.7	9.018.8
2004	R786-8567	Fault Shrimp	6080	2.340.34	12.843.0	4.014.4
2004	R788-87	Mat City	11540	10.141.10	13.543.8	9.842.3
2004	R788-85	Five Towers	190	2.340.17	16.843.7	11.344.0
2004	R788-CC	Five Towers	1044	0.210.02	14.343.3	10.344.0
2004	R789-85	Egg Drop Soup	156	2.340.62	18.344.8	5.515.5
2004	R790-856	Cave Vent	4380	11.311.37	12.842.4	3.311.9
2004	R791-856	Bacto Balls	477	4.010.19	16.445.2	2.011.0
2004	R791-87	Yellow Top	784	4.410.78	17.144.0	2.410.9
2004	R792-857	Champagne	3300	0.810.07	9.316.5	4.812.3
2004	R792-CC	Champagne	1350	0.510.03	6.443.9	3.512.3
2004	R793-81	Yellow Cone	218	2.010.24	15.444.6	5.512.3
2004	R793-857	Yellow Top	517	2.910.67	12.316.8	7.012.6
2004	R795-856	Fish Spa	7106	2.810.40	12.413.6	11.812.4
2006	J2-184-W	Fe-Mats	1486	2.910.28	10.414.3	5.912.5
2006	J2-184-B	Small Mat	5817	1.510.16	12.912.9	3.311.7
2006	J2-190-W	Fe-Mounds	3722	6.010.17	13.545.5	6.111.1
2006	J2-190-CC	Fe-Mounds	1631	6.310.50	11.548.3	6.110.0
2006	J2-191-W	Iceberg	617	0.410.15	6.012.2	7.912.6
2006	J2-197-W	Bubble Bath	1608	1.910.32	8.414.5	7.114.7
2006	J2-197-B	Fish Spa	5240	12.710.62	14.414.3	10.112.4

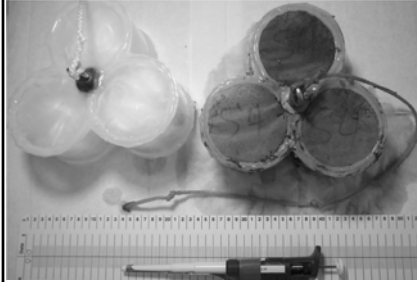
Difference is Bacteria →



SUMMARY

- Microbial community diversity is much greater along the Mariana Arc/Backarc than at either a hotspot volcano or at mid-ocean ridge along similar same spatial scales.
- This is most likely the result of the complexity of vent effluent chemistry and the ephemeral nature of hydrothermal venting along the Mariana Arc/Backarc.
- Cluster analysis of T-RFLP fingerprints reveals the microbial communities formed three distinct clusters designated Mariana Clusters I, II, and III.
- The Mariana Arc/Backarc hydrothermal vent systems support a Bacterial biodiversity hotspot which may be indicative of convergent plates boundary vent systems worldwide.

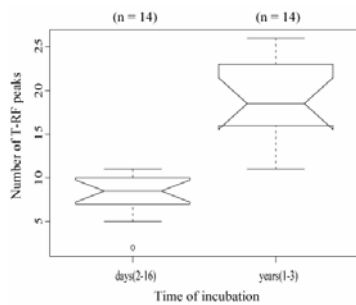
Microbial Growth Chambers (MGCs)



- Silica wool as inert substrate with $\sim 3\text{m}^2$ surface area
- Enclosed by 202 micron Nytex filter
- Can withstand 100°C
- Placed within flow for short- and long-term durations



Notched Box-and-Whisker Plot: Axial Seamount



- Entire dataset pooled into 2 bins
- Notches do NOT overlap indicating that the differences are highly significant
- Yields a first order approximation for bacterial colonization rates.

Quote: Baas Becking (1934)

(Referring to bacteria) - "everything is everywhere: but the milieu selects ... in nature and in the laboratory".

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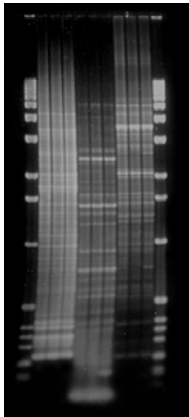
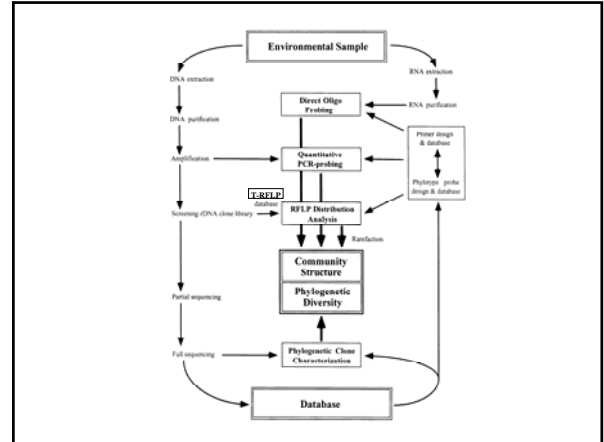


Andrea Curtis



Leslie Chao

**Submarine Ring of Fire
Diakoku Volcano 2006**



BOX, ERIC, & REP PCR

Lanes represent: Strains RL1, ES1, & ES2

- Minimal genetic variability among strains of Fe-oxidizing bacteria.
- Additional strains will determine if endemic or cosmopolitan at genomic level.