TABLE 4 Numbers and metabolic diversity of microorganisms in vent environments

| Samples | Number of microorganisms ${ }^{\text {a }}$ | Metabolic and/or phylogenetic groups |
| :---: | :---: | :---: |
| Diffuse-flow fluids $\left(2^{\circ} \mathrm{C}-80^{\circ} \mathrm{C}\right)$ | $10^{5}->10^{9} \mathrm{ml}^{-1}$; high numbers from Galapagos particles | Extremely high diversity of bacteria and archaea and include aerobes and anaerobes (see Table 1) |
| Smoker fluids $\left(250^{\circ} \mathrm{C}-400^{\circ} \mathrm{C}\right)$ | Not detected to $10^{7} \mathrm{ml}^{-1}$; high numbers correlate with evidence of phase separation ${ }^{\text {b }}$ | Hyperthermophilic methanogens and heterotrophic archaea isolated; evidence for other hyperthermophilic archaea and bacteria from molecular data ${ }^{\mathrm{c}}$ |
| Hydrothermal vent plume water $\left(2^{\circ} \mathrm{C}\right.$ in horizontal plume) | $\sim 10^{5}->10^{6} \mathrm{ml}^{-1}$ | $\mathrm{H}_{2^{-}}, \mathrm{CH}_{4^{-}}$, and $\mathrm{Mn}^{2+}$-oxidizing bacteria detected by activity measurements; Halomonas spp. isolated ${ }^{\text {d }}$ |
| Deep SW surrounding vents $\left(2^{\circ} \mathrm{C}\right)$ | $10^{3}-<10^{5} \mathrm{ml}^{-1}$ | Limited diversity of bacteria and archaea detected and enumerated using molecular methods |
| Sulfide structures | $>10^{8}$ per gram of sulfide on outer layers; $10^{5}$ per gram in interior | Outer layers have a mixture of archaea and bacteria and include metal oxidizers and methanogens; inner layers contain only archaea of unknown physiologies ${ }^{\text {e }}$ |
| Subsurface crust | Numbers are unknown | Different thermal groups of bacteria and archae detected from new eruptions; unique archaea isolated from subsurface fluids ${ }^{\mathrm{f}}$ |
| Microbial mats | $>10^{8}$ bacteria per gram | High numbers of sulfuroxidizing bacteria including Beggiatoa spp. and uncultured $\varepsilon$-proteobacteriag ${ }^{g}$ |
| Sediments | Surface of Guaymas sediments similar to microbial mats; numbers range from in sedimented ridges ${ }^{\text {h }}$ | Same as for microbial mats in surface layer with sulfatereducing bacteria and archaea dominating the deeper layers |

## TABLE 4 (Continued)

| Samples | Number of microorganisms ${ }^{\text {a }}$ | Metabolic and/or <br> phylogenetic groups |
| :--- | :--- | :--- |
| Animal <br> endosymbionts | $\sim 10^{10}$ per gram tissue | Sulfide-oxidizing bacteria most <br> common in hydrothermal vent <br> animals whereas methane-oxidizing <br> bacteria found at some cold seeps |
| Outer surfaces <br> of animals | Probably similar to <br> microbial mats | Methane oxidizers found on surface <br> shells of some limpets and worm <br> tubes |
|  |  | filamentous metal-oxidizing <br> bacteria found on rear surfaces <br> of Alvinella worms |

${ }^{\text {a }}$ Numbers usually determined by epifluorescence microscopy or quantitative lipid analyses.
${ }^{\text {b}}$ Baross \& Deming 1995.
${ }^{\circ}$ Takai et al. 2000.
${ }^{\text {d }}$ Cowen et al. 1986, 1998; Lilley et al. 1995; Kaye \& Baross 2000.
${ }^{\text {e }}$ Harmsen et al. 1997a, Schrenk et al. 1999a.
fdelaney et al. 1998; Summit \& Baross 1998, 2001.
\&Jannasch 1995, Longnecker \& Reysenbach 2001, Nelson et al. 1989, Moyer et al. 1995.
${ }^{\text {h }}$ Parkes et al. 1994, Summit et al. 2000.
${ }^{i}$ De Angelis et al. 1991.
${ }^{\mathrm{j}}$ Cary \& Stein 1998.

