crust, just because he thought it was interesting. Over lunch one day, he heard from colleagues that engineers were planning to bury the Alaska oil pipeline in permafrost. Lachenbruch realized, when no one else seemed to, that the heat from fluid friction within the pipe would surely melt the permafrost and wreak disasters on Alaska. The seemingly irrelevant researcher quickly convinced the engineers, and the pipeline was elevated above the permafrost.

People like Lachenbruch "see problems in ways other people can't," says Dalrymple, and are thus a crucial component of an organization like the USGS. "Even though you're downsizing, there has to be some critical mass [of basic research] you maintain. I

don't know what it ought to be, but it probably has to be larger than 5 or 10%."

Eaton disagrees. In a reply to concerns expressed in a letter from Barton, Eaton argued that "the 10 to 12% of [Water Resources Division] staff who do basic research (as opposed to data-gathering and analysis) strike a balance that, if applied to [the Geologic Division], would provide more than ample room for" work like Lachenbruch's. And some staffers see the paring of research in many parts of the Geologic Division as inevitable. One scientist who has shuttled between research and management at headquarters argues that it is proper for the Geologic Division to focus more on understanding local processes, such as volcanic

eruptions or landslides, and less on basic research and the broad geologic mapping that some traditionalists still view as the core of the division.

"We're not a basic research organization," says the staffer. "We have to show more impact on society, policy, or economics. Some have come to feel they can do science for science's sake and have lost touch with the underlying rationale of why the tax-payers should pay their salary." Applied science and gathering fundamental knowledge "are not separate," he notes; "it's a matter of balance." Finding the right balance, and retaining the right people to make it work, will be the challenge.

-Richard A. Kerr

GENETICS RESEARCH_

NIH's "Gay Gene" Study Questioned

Two years ago, geneticist Dean Hamer of the National Cancer Institute published a study of 40 pairs of brothers—all gay—reporting that their sexual orientation was influenced by their genes. Family pedigree data indicated that the men had inherited a factor for gayness from their mothers. Hamer and his team zeroed in on the X chromosome (passed to males only by their mothers), scanning it for genetic markers that the gay men might have in common. Based on these linkage studies, Hamer concluded he had found a gay genetic factor at the tip of the X chromosome (*Science*, 16 July 1993, pp. 291 and 321).

This report—offering the first molecular evidence that human sexual orientation might be determined genetically-sparked controversy, and lots of it. But in recent months Hamer's work has begun to face more serious technical questions—one in a confidential setting, the other in public. The confidential investigation is being carried out by the Office of Research Integrity (ORI) in the Department of Health and Human Services. And in a more public forum, the research is also being questioned by George Ebers, a neurogenetics researcher at the University of Western Ontario in London, Ontario, who is trying to confirm Hamer's result. Although Ebers says his research is similar to that of the National Institutes of Health (NIH) team, so far, he told a gathering at Cold Spring Harbor Laboratory in March, he hasn't replicated Hamer's finding.

News of the ORI inquiry broke on 25 June when the Chicago Tribune reported that a former junior member of Hamer's lab at the NIH had raised questions about Hamer's research. According to the Tribune, the postdoc, a co-author with Hamer who did gene mapping studies in his lab, triggered the ORI inquiry in March 1994 when she challenged unspecified methods of data selec-

tion. NIH declines to comment. Lyle Bivens, ORI's director, says his office only conducts investigations at NIH after NIH has completed an inquiry, but he didn't discuss this case: "We cannot confirm or deny the existence of any investigation."

Hamer also declines comment. But two geneticists contacted by *Science*, both intramural researchers at NIH, confirm that ORI is investigating Hamer's work. Both de-



Drawing fire. Dean Hamer, leader of the NIH team that found a locus for gay behavior on the X chromosome.

manded anonymity. ORI is looking into allegations that Hamer "selectively reported his data," according to the Tribune. Another question that may have prompted an inquiry, says an NIH researcher, is: How did Hamer select subjects? There is little consensus within the scientific community on the best way to identify gay members of a family; women, for example, may be more reliable informants than men. It is not clear whether ORI's investigation focuses on Hamer's 1993 paper or on a follow-up study on a new group of 33 pairs of gay brothers (Science, 16 June, p. 1571). One of Hamer's co-authors on the second paper, geneticist Stacey Cherny of the University of Colorado, Boulder, says this still-unpublished research supports Hamer's original finding, but with a lower level of statistical significance.

As the confidential ORI investigation moves forward, a general debate on Hamer's findings is taking place in public, sparked by Ebers and George Rice, collaborators at the University of Western Ontario. Ebers says he agrees with Hamer's view that gay behavior is probably inherited, but sees no reason to focus linkage studies on the X chromosome. About 4 years ago, Ebers says, he began to

look into the genetics of gay behavior "as a matter of personal curiosity." With Rice's help, he collected data on more than 40 pairs of gay brothers—the number Hamer studied. But unlike Hamer, Ebers found no evidence that gayness is passed from mother to son—"not even a trend in favor of X-linkage."

Hamer, in an e-mail response to *Science*, says Ebers' "research design is very different than our own and cannot be interpreted to either refute or confirm our findings." Ebers and Rice may have missed the X-linkage, according to Hamer, because "they made no

effort to select families that display the maternal pattern of inheritance," and thus diluted the critical genetic information in a sea of noise. Hamer notes that different groups studying complex genetic traits often reach different conclusions.

Ebers doesn't think that his selection of subjects biased the conclusions. But he does sympathize with Hamer's discomfort at the criticism he's getting, and he notes that the whole matter should be regarded as "an open question," requiring "a lot more work to sort it out." Hamer, for his own part, doubts there would be a fuss "if we were working on any topic other than homosexuality."

-Eliot Marshall