NEWS & COMMENT

trol goes out of whack for a while," says Markham. He plans to study how Merbold's eye control changes as he adapts to weightlessness, and he hopes the information will help in screening astronauts for their susceptibility to space motion sickness.

For his experiment, Markham will use an instrumented helmet brought up for the 1992 German mission to Mir. The device records eye movements on video while sensors record the actual movements of the astronaut's head. Clarke in Berlin will use the helmet for a more fundamental study of the vestibular system: Motion sensors in the inner ear are influenced by gravity, so once it is removed, it will be possible to see directly how a simple movement of the head, say tilting it to one side, is detected by the sensor and so directs eye movement. The long duration of the flight "takes the pressure off the experiment," he says. Experiments like Clarke's are designed to shed light not only on the problems caused by prolonged weightlessness but also on how the human body functions on Earth. "We are using space as a tool. It is no different from ground-based research," says Oser. The distribution of blood and other fluids throughout the body, for example, is dominated by hydrostatic pressure caused by gravity. By removing gravity, scientists can see how volume shifts are regulated. "Space unmasks the mechanism," says Oser.

In one such study of the control systems of bodily fluids, devised by Peter Bie of the University of Copenhagen, Denmark, Merbold will draw some of his own blood with the help of a Russian colleague, mix it with salt solution, and reinfuse it. By comparing blood samples taken during the flight as well as just before and after it, Bie hopes to learn how salt defuses through the body. Other

_____ cDNA SEQUENCES ___

HGS Opens Its Databanks—For a Price

 ${f T}$ he biologist-entrepreneurs who run the world's largest commercial human gene sequencing project are announcing a plan this week to share their data with academic researchers-at a price. Human Genome Sciences (HGS) and its nonprofit research partner, the Institute of Genomic Research (TIGR) of Rockville, Maryland, say they're ready to share detailed information they've collected on 35,000 complementary DNAs representing genes of unknown function that are expressed in human tissue. But those who use HGS's proprietary data would first have to agree to give HGS the option of licensing derivative products at a reasonable royalty rate. And they would have to provide 30 to 60 days' prior notice of publication.

William Haseltine, president of HGS, says his company will soon give researchers broad access to genetic data on these terms after HGS and TIGR have published a paper this fall describing the cDNAs they've analyzed. As *Science* went to press, Haseltine was preparing to discuss the plan at a meeting of the international Human Genome Organization in Washington, D.C., attended by the major players in the human genome effort. One of the players, Maxwell Cowan, chief scientific officer of the Howard Hughes Medical Institute, said he thought the group would find the terms "acceptable."

Even if Cowan's prediction is correct, however, HGS may soon may face some new competition from a venture planning to make similar information available with no strings attached. Last week, Merck & Co., the pharmaceutical giant, announced that it will bankroll a new effort based at Washington University in St. Louis to sequence human cDNAs. Unlike HGS, Merck plans to put sequence data into a public database as soon as they become available. According to a Merck executive, the company hopes to encourage basic research that will go beyond sequencing genes and begin to explain their function. Merck officials say they want to speed up the discovery of new drugs.

The first big cDNA drive began in 1992, when HGS and TIGR launched an effort to clone and patent human cDNA using robotic methods developed by J. Craig Venter, a scientist who left the National Institutes of Health (NIH) in 1992 to take charge of TIGR. In 1993, Haseltine and Venter signed an agreement with SmithKline Beecham (SB) to develop medical products, and since then, their sequencing project has received \$85 million from SB and a promise of roughly \$40 million more. So far, according to Haseltine, the partnership has generated 120 million expressed nucleotides, identified more than 80,000 unique sequences, validated the process by independently confirming the existence of 202 of the 248 human genes already named in reports, and filed for patents on 70 full-length cDNAs.

The companies are now ready to share this "magnificent achievement," Haseltine says. In the past, Haseltine notes, the HGS-TIGR partnership has shared data in tailored agreements on a "one-off basis" with individual scientists. For example, a collaboration with Bert Vogelstein of Johns Hopkins University led to the discovery of a colon cancer gene. But in recent months, HGS, TIGR, and SB have drafted a new "material transfer agreement" to make it easier for scientists to tap into their data.

The goal of the new arrangement, says Haseltine, is to speed up the process of ge-

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experiments into bone and muscle loss will be limited to examinations of Merbold shortly before and after the flight because of the lack of suitable instruments onboard.

If all goes according to plan, ESA will have another astronaut on Mir in a year's time, and some of these experiments will be repeated then. The extra time until Euromir-95 will allow ESA to complete development of a bone densitometer that will use ultrasound to monitor wasting of the astronaut's heel bone during the mission. ESA also hopes to send up equipment to analyze respiration and urine during the flight, lessening the need to bring samples back to Earth. By the time the international space station is up and running around the year 2000, the Europeans and Russians will already have amassed substantial data on space physiology—and at a bargain price.

-Daniel Clery

netic research. The gesture may also give HGS and TIGR a public relations boost. In recent months they've been criticized for trying to hold valuable scientific information too close to the chest. Some researchers have balked at restrictions HGS has placed on commercial use of the data in the past. For example, *Science* has learned that the Baylor College of Medicine, the Howard Hughes Medical Institute, and NIH objected to the conditions of an HGS data-sharing agreement proposed earlier this year.

And that's one reason why Merck jumped into the sequencing business. Merck feels that "this should be public information," says Alan Williamson, Merck's vice president for global research strategy. The real "value added," as Williamson sees it, is in the basic biology that must still be done to determine the function of the DNA sequences. Some HGS and TIGR officials, however, regard Merck's move as a "spoiler strategy" designed to "throw a wrench in the machinery."

Details of Merck's project are still being negotiated. But Williamson and Robert Waterston, director of Washington University's genetic sequencing laboratory, say the initial Merck grant will amount to several million dollars and last for 18 months or, as Williamson says, long enough to get the job done. Waterston says he expects the new project will be up and running before the end of the year. He adds that he's delighted by the recognition of the lab's accomplishments, especially Richard Wilson's effort to sequence DNA from a model organism, the nematode Caenorhabditis elegans. While the "public sometimes has trouble understanding why NIH would spend money on sequencing a nematode," Waterston says, "Merck didn't have any trouble."

–Eliot Marshall