

Growing Joints Use Their Noggins

Some arthritis sufferers might wish their joints would just go away, but new research presented on page 1455 of this issue shows that jointlessness is not a happy alternative. Molecular embryologists Richard Harland and Lisa Brunet at the University of California, Berkeley, and Jill and Andrew McMahon at Harvard University, have found that mice lacking *noggin*, a gene first discovered as important in brain and nerve development, have no joints at all. Instead, they have stubby, continuous limbs—along with a fatal array of other developmental defects.

noggin is “a new link in the chain of creation of limbs” and the rest of the skeleton, says cell biologist Bjorn Olsen of Harvard Medical and Dental Schools. Harland’s finding is a step toward a more detailed understanding of embryonic development, adds molecular biologist Sejin Lee of Johns Hopkins University, and offers potential medical benefits in diseases where there is too much bone or even too little.

The new finding is just the latest role for *noggin*, which William Smith, then in Harland’s lab, and Harland identified in 1992 after setting out to find the “neural inducer,” a molecule that orders cells to become brain and nervous system tissue in early embryos. The gene earned its name when they found that frog embryos injected with its messenger RNA grew exceptionally large heads. The *noggin* protein also mimics the activity of a powerful piece of tissue in the developing frog known as “Spemann’s organizer,” which can make back-of-the-body (dorsal) tissue out of front-side (ventral) tissue. Finally, 2 years ago, Harland’s team showed that in binding assays and cell culture, *noggin* inhibits powerful proteins that stimulate bone growth, the so-called bone morphogenetic proteins (BMPs).

With *noggin* playing all these developmental roles, Harland half expected that when the team turned off the gene in their mice, the resulting animals would be no more than “a ball of mush.” Indeed, the knockout mice did not survive until birth. But they developed enough to offer a new insight into *noggin*’s function.

The mouse showed a variety of intriguing skeletal abnormalities. “Every single bone is affected,” says Harland, with the most obvious defects, such as shorter bones, in the verte-

brae, ribs, and limbs. And in keeping with the gene’s role in the brain, “there are very clearly characterized neural defects,” including occasionally a brain and spinal cord not enclosed by bone. But the bones in the heads and upper bodies of the mice are much less affected by the knockout than bones farther toward the animals’ tails. And dorsoventral patterning doesn’t seem to be much affected. This implies that *noggin* has counterparts that can perform its functions near the head and in dorsoventral patterning, says Harland.



Dis-jointed. Mice missing the *noggin* gene have paws lacking joints.

Most strikingly, the mice appear to lack all joints. Instead, their limbs are nearly continuous segments of bone, flanked by excess cartilage. That makes sense, says Harland, because during normal development, cartilage is laid out like a pencil sketch in the shape of the bones-to-be. Bone gradually fills in this cartilaginous sketch except in predetermined locations such as at the ends of the putative bones. In those spaces, the cartilage then disappears, leaving room for joints like knees and knuckles. In the knockout mice, the cartilage does not

do this disappearing act. Without *noggin*, the thinking goes, bone-forming proteins go out of control, recruiting additional cells from neighboring areas into the prebone cartilage.

But here too *noggin* apparently does not act

alone, according to work by Harland’s and other labs. Instead, it apparently sends signals to a BMP family member called GDF-5, which has been shown to be important in joint formation, and also interacts with another limb-building protein, Sonic hedgehog. Therefore, says Olsen, *noggin* “is not a master molecule” that regulates everything else. Because BMPs are regulated by the powerful family of patterning genes known as *hox*, he proposes that *noggin* is a link in the pathway, somewhere downstream of *hox* genes and upstream of BMPs, that governs the patterning of limbs.

Uncovering the molecular basis of this pathway has clinical implications, because many diseases, from osteoarthritis to osteoporosis, involve either too much bone or too little. Biotech companies are already avidly testing BMPs as potential drugs; two are in clinical trials now for healing bone breaks. But these molecules are, if anything, too powerful, says Lee. “The challenge,” he says, “is to limit bone growth to what is clinically desirable.”

Limiting bone growth is where *noggin* might come in. Regeneron Pharmaceuticals of Tarrytown, New York, is studying whether *noggin* and other BMP inhibitors can put the brakes on the excess bone growth that arises in about 10% of hip replacement patients as well as in some patients with osteosarcomas and prostate cancer metastases, says biochemist Neil Stahl of Regeneron.

Systemic use of *noggin* could have unwanted side effects, because the BMPs it inhibits are found everywhere from skin to gut to bone, warns Rik Derynck, a cell and developmental biologist at the University of California, San Francisco. But eventually, drug companies might use their noggins to provide novel treatments for overgrown bone.

—Steven Dickman

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INFECTIOUS DISEASES

New Method Churns Out TB Mutants

By and large, bacteria are much easier to study in the laboratory than more complex, multicellular organisms. But every rule has its exceptions, and for microbiologists one of the cruellest has been *Mycobacterium tuberculosis*, the pathogen that causes tuberculosis (TB). The microbe’s recalcitrance in the lab has hindered researchers in their efforts to design better drugs for combating TB, which is the world’s leading killer among infectious diseases, claiming more than 3 million lives worldwide every year. Now, that impasse may be at an end.

In order to ferret out new drug targets, researchers want to identify the genes that pathogens need to survive and infect the host—a task usually accomplished by creating wholesale mutations in the microbial genome and then

screening for mutants defective in those abilities. Until a few months ago, this was very hard to do with *M. tuberculosis*, partly because the vehicles typically used to create the mutations—small bits of DNA called transposons that insert randomly into the genome and inactivate any gene they happen to interrupt—do not readily penetrate the microbe’s tough, waxy coat.

Last fall, however, a team of scientists led by microbiologist William Jacobs and immunologist Barry Bloom of the Albert Einstein College of Medicine in New York City, reported creating a new kind of vehicle—a cross between a bacterial virus and a circle of DNA called a plasmid—that’s much more efficient at producing muta-