

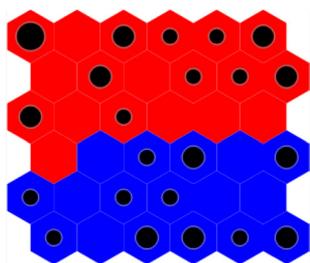
- 15527 Genomes reveal bacterial lifestyles
 15538 Glowing markers of cell death in bananas
 15583 Media multitaskers pay a cognitive price
 15599 Red bioluminescent protein detects tumors in live animals

- 15714 Cultured hepatocytes aid in drug screening
 15720 Using fat to generate stem cells
 15768 Insulin-producing β cells created from patients with Type 1 diabetes
 15837 Free heme may contribute to severe malaria

ENVIRONMENTAL SCIENCES

Genomes reveal bacterial lifestyles

Researchers have used genomic sequence comparisons to predict an ocean bacterium's trophic strategies, which may provide a technique to analyze the diversity of ocean bacteria,



Self-organizing map showing two clusters of organisms.

many of which are difficult to culture in the lab. Federico Lauro et al. compared the genomes of two common ocean bacteria: the copiotrophic *Photobacterium angustum*, which lives in warm, nutrient-rich waters and is fast to grow and divide, and the oligotrophic *Sphingopyxis alaskensis*, which lives in nutrient-poor, Arctic waters and grows more slowly. The authors

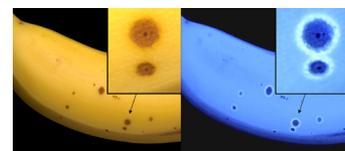
analyzed the number, patterns, and variations in gene families. In one family, copiotrophic bacteria had many selective transporter proteins to quickly absorb plentiful nutrients, whereas oligotrophic bacteria had a smaller number of highly efficient transporter proteins to extract what little nutrition is available in Arctic oceans. The authors found differences in other genes related to nutrient and energy usage, reflecting the bacteria's adaptation to their environments. The authors used their technique to analyze 124 other oceanic bacteria, and found that oligotrophic bacteria outnumbered copiotrophic bacteria in worldwide samples of ocean water, despite the fact that copiotrophic bacteria are easier to culture in the lab. — P.D.

"The genomic basis of trophic strategy in marine bacteria" by Federico M. Lauro, Diane McDougald, Torsten Thomas, Timothy J. Williams, Suhelen Egan, Scott Rice, Matthew Z. DeMaere, Lily Ting, Haluk Ertan, Justin Johnson, Steven Ferreira, Alla Lapidus, Iain Anderson, Nikos Kyrpides, A. Christine Munk, Chris Detter, Cliff S. Han, Mark V. Brown, Frank T. Robb, Staffan Kjelleberg, and Ricardo Cavicchioli (see pages 15527–15533)

CHEMISTRY, PLANT BIOLOGY

Glowing markers of cell death in bananas

As a banana transitions from ripe to rotten, the yellow peel develops brown and black spots that grow rapidly, eventually enveloping the whole fruit. The spots are a sign of local cell death and reveal necrotic tissue. Simone Moser et al. found that aging bananas glow blue under UV light, and the black spots are surrounded by bright blue



Ripening bananas exhibit luminescent blue halos.

halos. The fluorescent blue color is caused by the breakdown of chlorophyll into fluorescent chlorophyll catabolites (FCCs). In bananas, the authors show that FCCs accumulate and are persistent molecules that reveal a previously unknown pathway for chlorophyll breakdown and may prove particularly useful to researchers as noninvasive markers of cell death. The authors note that nonfluorescent chlorophyll catabolites are antioxidants and may extend the viability of aging tissues, although it is not known whether FCCs would do the same for bananas. Whereas the biological role of these fluorescent molecules is unclear, the authors speculate that fruit-eating animals may detect the fluorescent halos as a sign of ripe food.

— B.P.T.

"Fluorescent chlorophyll catabolites in bananas light up blue halos of cell death" by Simone Moser, Thomas Müller, Andreas Holzinger, Cornelius Lütz, Steffen Jockusch, Nicholas J. Turro, and Bernhard Kräutler (see pages 15538–15543)

PSYCHOLOGICAL AND COGNITIVE SCIENCES

Media multitaskers pay a cognitive price

Media multitasking is becoming ubiquitous as technology proliferates and people are able to simultaneously consume multiple forms of media, including music, e-mail, video, and other applications. How the human brain copes with this shift is unclear.

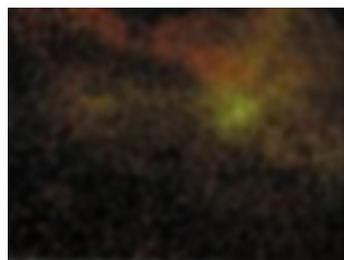
ties. The authors isolated adipose cells from adults between the ages of 45 and 60 and attempted to reprogram the cells into stem cells using an established genetic targeting method. At the same time, they began the same procedure with adult skin cells. The adipose cells produced adult stem cells twice as fast and 20 times more efficiently than skin cells. In addition, the fat cells did not require addition of mouse feeder cells, which could complicate the method's future use, according to the authors. — T.H.D.

“Feeder-free derivation of induced pluripotent stem cells from adult human adipose stem cells” by Ning Sun, Nicholas J. Panetta, Deepak M. Gupta, Kitchener D. Wilson, Andrew Lee, Fangjun Jia, Shijun Hu, Athena M. Cherry, Robert C. Robbins, Michael T. Longaker, and Joseph C. Wu (see pages 15720–15725)

DEVELOPMENTAL BIOLOGY

Insulin-producing β cells created from patients with Type 1 diabetes

Type 1 diabetes, also known as juvenile diabetes, is an autoimmune disease that destroys the insulin-producing β cells in the pancreas. The molecular or genetic defects that trigger



Differentiation of human pluripotent stem cells to β -like cells.

T1D are unknown. Animal models have been useful, but it is unclear to what extent they replicate the human disease. René Maehr et al. developed a tool for studying T1D that may also work as a cell replacement therapy. Using three transcription factors—OCT4, SOX2, and KLF4—the authors reprogrammed adult fibroblast cells taken from T1D patients and transformed them into induced pluripotent stem cells (iPS). These T1D iPS cells are a genetic match to the patient from which they are derived and have the potential to develop into many cell types in the body. The authors coaxed the T1D iPS cells to become insulin-producing β cells—though the process was not very efficient. Still, the work provides proof of principle that genetically matched, insulin-producing cells can be made from patients with T1D. If coupled with a method to block rejection, these cells could be useful for transplantation, but their

principal value is as a tool to study how and why diabetes occurs, according to the authors. — B.P.T.

“Generation of pluripotent stem cells from patients with type 1 diabetes” by René Maehr, Shuibing Chen, Melinda Snitow, Thomas Ludwig, Lisa Yagasaki, Robin Goland, Rudolph L. Leibel, and Douglas A. Melton (see pages 15768–15773)

MEDICAL SCIENCES

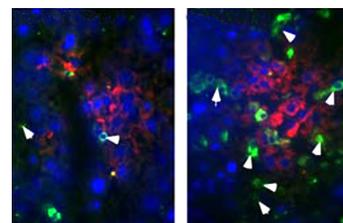
Free heme may contribute to severe malaria

Malaria remains one of the world's deadliest diseases, responsible for up to 3 million deaths annually. The causative parasites, which are from the *Plasmodium* family of protozoans, infect and replicate in red blood cells. When the pathogens emerge, they rupture these cells, and hemoglobin and other normally sequestered molecules enter the blood

stream where they can be oxidized and produce deleterious chemicals such as free heme, hemoglobin's iron center. Elsa Seixas et al. report that free heme's pro-oxidant properties

can lead to the death of *Plasmodium*-infected mice. The survival of an infected host relied on the heme-catabolizing enzyme, heme oxygenase-1 (HO-1), which prevented the cytotoxic effects of free heme; *plasmodium*-infected mice that lacked the enzyme developed a lethal form of hepatic failure. Although HO-1 did not inhibit the protozoan's ability to replicate, the enzyme prevented free heme from provoking TNF-mediated programmed cell death in hepatocytes, affording so-called host “tolerance” against *Plasmodium* infection. A pharmacological antioxidant, N-acetylcysteine (NAC), mimicked protective effect of HO-1 and suppressed the development of liver failure in infected mice. This research may lead to therapeutics that promote host tolerance against *Plasmodium* infection and could reduce the protozoan's ability to cause severe forms of malaria, according to the authors. — F.A.

“Heme oxygenase-1 affords protection against noncerebral forms of severe malaria” by Elsa Seixas, Raffaella Gozzelino, Angelo Chora, Ana Ferreira, Gabriela Silva, Rasmus Larsen, Sofia Rebelo, Carmen Penido, Neal R. Smith, Antonio Coutinho, and Miguel P. Soares (see pages 15837–15842)



Liver of noninfected (Left) or Pcc-infected (Right) mice.