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ABOVE: ROGERS FUND, 1930, CHARLES K WILKINSON

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Archaeobotanists at UCL helped Mott and a team of collaborators choose a handful of well-preserved husks from the museum's collection of ancient emmer wheat, a variety native to the Near East and one of the first crops to be domesticated in the region, from which the researchers selected two husks for DNA extraction. After carefully removing the husks from the box, photographing them, and wrapping them in foil, the scientists transported the centuries-old plant material to a freshly bleached cleanroom used exclusively to process ancient and forensic samples. There, team member Laura Botigué, a population geneticist and visiting researcher from the Centre for Research in Agricultural Genomics (CRAG) in Barcelona, Spain, donned a hairnet, two Tyvek suits, two pairs of latex gloves, and a mask—part of a protocol designed to avoid contaminating the samples with her own cells. Uncertain how the delicate husks would hold up to the standard decontamination protocol of bleaching the samples, Botigué bleached one and left the second untouched. Then, to lyse the plant's cells, she put the samples in a rotator that gently shook the husks inside an oven over the next several days. Finally, she used a centrifugation protocol to separate any DNA from the degraded cell walls and proteins.

Once the samples had been prepped and delivered to the UCL Genomics facility for sequencing, it was a waiting game to see if the procedure had yielded any readable wheat DNA. “This is the more stressful part,” Botigué says. Because they lack the type of protective collagen matrix found in bones, plants don’t preserve ancient DNA as well as animals. “You finish, the DNA is theoretically extracted, but you don’t see it in the tube,” says Botigué. “You’re in the blind until you hear back from the sequencing services.”

Within just a few weeks, the team got good news. For the husk that Botigué had bleached, about two-thirds of the reads aligned with genomes of modern wild and domesticated emmer wheat varieties—a relatively good success rate for ancient DNA, according to evolutionary geneticist Michael Scott, a postdoc in Mott’s lab who conducted the bioinformatics analysis of the sequences. “The first surprise was how well it worked,” he says. “It appears that the dry conditions in Egypt were good for DNA preservation.” The unbleached husk had yielded a smaller quantity of sequences, but those fragments mostly matched the ones in the bleached sample, validating the identity of those sequences as coming from the ancient wheat samples rather than from contaminants.

The museum wheat, which carbon dating showed was from between 1130 and 1000 BC, was genetically much more similar to modern domesticated varieties than to modern wild ones, suggesting that the plant lineage the samples came from had already been domesticated. Specifically, the sequences most resembled those of modern domesticated strains grown in Turkey, Oman, and India. There was also evidence for genetic exchange between the museum wheat strain and the wild emmer wheat that grew in the Levant, a large region in the Eastern Mediterranean that was a center of agricultural development in the Neolithic period, and where emmer was first cultivated. The genetic exchange could have occurred before the wheat’s introduction to Egypt from the Levant, says Scott. Alternatively, it’s possible that the ancient Egyptians’ wheat was able to interbreed with wild wheat in the Southern Levant thanks to interactions between the people in the

two regions.

“With big data and with a really good analysis method they were able to detect this gene flow,” says M. Timothy Rabanus-Wallace, an agricultural geneticist at the Leibniz Institute of Plant Genetics and Crop Plant Research in Germany who coauthored a perspective published alongside the study in *Nature Plants* last October. “It’s fascinating to see this gene flow happening . . . in an area important for human history.”

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The bioinformatics analysis also uncovered some genetic variants in the ancient samples that weren’t found in any of the modern emmer wheat genomes the researchers studied. If these variants helped the wheat survive in arid locations around the Near East, perhaps introducing those sequences into modern varieties could help make them more sustainable or more drought resistant, Scott says, though he admits that this “is very much just an idea.”

The detection of ancient genetic variation is a notable achievement because wheat genomes are large—three to five times the length of a human genome—and repetitive, making the “analysis . . . incredibly complex,” says James Breen, head of the bioinformatics core at the South Australian Health and Medical Research Institute who reviewed the study and coauthored the perspective with Rabanus-Wallace, a PhD student in his lab at the Australian Centre for Ancient DNA at the time. “So being able to find unique pieces of DNA in that genome is very difficult.” He adds that after a couple of additional validation tests performed by the UCL team, he was convinced that “the data that came out was legitimately ancient.”

Botigué and Scott emphasize that the study is primarily a proof of concept that museum-kept plant samples can yield readable genetic material. “We were able to look at DNA from specimens that had been stored in the museum for over 90 years without special preservation conditions—the museum was actually even bombed and flooded during wartime,” says Scott. “We think our study helps

demonstrate the importance of museum collections as sources of genetic data, which”—in combination with new samples—“can be used to uncover the history of selection on crops and their movement around the globe.”

See “The Narluga: New Insights from Old Bones”

“I think that’s one of the biggest values of ancient DNA in plants,” adds Nathan Wales, an archaeologist at the University of York who was not involved in Scott and Botigué’s study—“to draw connections between different cultures and the different agricultural products they were growing and trading, and seeing how that changed over time.”