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appear to be essential for anhydrobiosis. Attention is now focusing on various 'omic' approaches to the study of anhydrobiosis [5] to explore whether there is a 'desiccome' — a set of genes associated with desiccation survival and anhydrobiosis [6]. Of particular interest is desiccation-induced protein synthesis and the role of *lea (late embryogenesis abundant*) genes [19].

So will C. elegans dauers prove a useful model for the study of anhydrobiosis? Undoubtedly, but with a note of caution. The desiccation survival abilities of dauer-constitutive mutants are modest with 10% survival after exposure to 0% RH, following preconditioning at 98% RH. The survival of wild-type dauers is even lower [10]. Nematodes have a range of desiccation survival abilities [14], even within the same genus [20]: some will survive direct exposure to 0% RH with high levels of survival ('innate dehydration strategists' [14]). Perhaps our understanding of anhydrobiosis will be advanced by comparing the biology of organisms with different abilities to survive desiccation and anhvdrobiosis.

References

1. Watanabe, M. (2006). Anhydrobiosis in invertebrates. Appl. Entomol. Zool. 41, 15–31.

- Clegg, J.S. (2001). Cryptobiosis a peculiar state of biological organization. Comp. Biochem. Physiol. B-Biochem. Mol. Biol. 128, 613–624.
- Reardon, W., Chakrabortee, S., Pereira, T.C., Tyson, T., Banton, M.C., Dolan, K.M., Culleton, B.A., Wise, M.J., Burnell, A.M., and Tunnacliffe, A. (2010). Expression profiling and cross-species RNA interference (RNAi) of desiccation-induced transcripts in the anhydrobiotic nematode *Aphelenchus avenae*. BMC Mol. Biol. *11*, 6.
- Wharton, D.A., Petrone, L., Duncan, A., and McQuillan, A.J. (2008). A surface lipid may control the permeability slump associated with entry into anhydrobiosis in the plant parasitic nematode *Ditylenchus dipsaci*. J. Exp. Biol. 211, 2901–2908.
- Schill, R.O., Mali, B., Dandekar, T., Schnolzer, M., Reuter, D., and Frohme, M. (2009). Molecular mechanisms of tolerance in tardigrades: New perspectives for preservation and stabilization of biological material. Biotechnol. Adv. 27. 348–352.
- Biotechnol. Adv. 27, 348–352.
 Boschetti, C., Pouchkina-Stantcheva, N., Hoffmann, P., and Tunnacliffe, A. (2011). Foreign genes and novel hydrophilic protein genes participate in the desiccation response of the bdelloid rotifer *Adineta ricciae*. J. Exp. Biol. 21, 59–68.
- Mitsumasu, K., Kanamori, Y., Fujita, M., Iwata, K., Tanaka, D., Kikuta, S., Watanabe, M., Cornette, R., Okuda, T., and Kikawada, T. (2010). Enzymatic control of anhydrobiosisrelated accumulation of trehalose in the sleeping chironomid, *Polypedilum* vanderplanki. FEBS J. 277, 4215–4228.
- 8. Maher, B. (2009). Biology's next top model? Nature 458, 695–698.
- 9. Blaxter, M. (2011). Nematodes: The worm and its relatives. PLoS Biol. 9, e1001050.
- Erkut, C., Penkov, S., Khesbak, H., Vorkel, D., J.-M., V., Fahmy, K., and Kurzchalia, T.V. (2011). Trehalose renders the dauer larva of *Caenorhabditis elegans* resistant to extreme desiccation. Curr. Biol. *21*, 1331–1336.
- 11. Grant, W., and Viney, M. (2011). The dauer phenomenon. In Molecular and Physiological

Basis of Nematode Survival, R.N. Perry and D.A. Wharton, eds. (Wallingford: CABI Publishing), pp. 99–125.

- Barrière, A., and Félix, M.A. (2007). Temporal dynamics and linkage disequilibrium in natural *Caenorhabditis elegans* populations. Genetics 176, 999-1011.
- Fitch, D.H.A. (2005). Evolution: An ecological context for C. elegans. Curr. Biol. 15, R655–R658.

 Perry, R.N., and Moens, M. (2011). Survival of parasitic nematodes outside the host. In Molecular and Physiological Basis of Nematode Survival, R.N. Perry and D.A. Wharton, eds. (Wallingford: CABI Publishing), pp. 1–27.

- Crowe, J.H., Hoekstra, F., and Crowe, L.M. (1992). Anhydrobiosis. Annu. Rev. Physiol. 54, 579–599.
- Ratnakumar, S., and Tunnacliffe, A. (2006). Intracellular trehalose is neither necessary nor sufficient for desiccation tolerance in yeast. FEMS Yeast Res. 6, 902–913.
- Lapinski, J., and Tunnacliffe, A. (2003). Anhydrobiosis without trehalose in bdelloid rotifers. FEBS Lett. 553, 387–390.
- Womersley, C.Z., and Higa, L.M. (1998). Trehalose - its role in the anhydrobiotic survival of *Ditylenchus myceliophagus*. Nematologica 44, 269–291.
- Tunnacliffe, A., and Wise, M. (2007). The continuing conundrum of the LEA proteins. Naturwissenschaften 94, 791–812.
- Shannon, A.J., Browne, J.A., Boyd, J., Fitzpatrick, D.A., and Burnell, A.M. (2005). The anhydrobiotic potential and molecular phylogenetics of species and strains of *Panagrolaimus* (Nematoda, Panagrolaimidae). J. Exp. Biol. 208, 2433–2445.

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Associative Memory: Without a Trace

Some transient sensory stimuli can cause prolonged activity in the brain. Trace conditioning experiments can reveal the time over which these lasting representations can be utilized and where they reside.

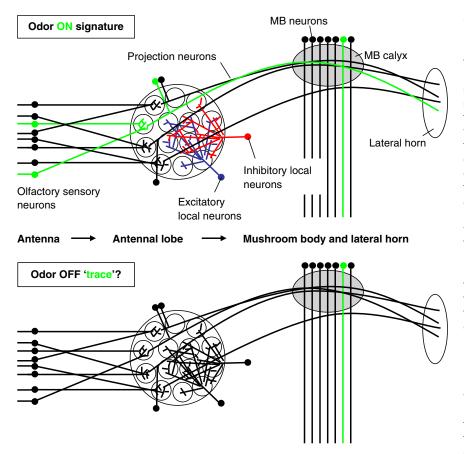
Emmanuel Perisse and Scott Waddell

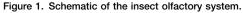
Associative learning allows an animal to know that the presence of a stimulus - the conditioned stimulus or CS, such as a smell or taste - predicts another is forthcoming - the unconditioned stimulus or US, a reward or punishment. Since the classic experiments of Pavlov [1], we have known that associative conditioning is most efficient when the CS and US presentations overlap in time (delay conditioning). However, learning can also occur when there is a pronounced gap between the CS and US (trace conditioning). Perhaps the most

extreme case is in conditioned taste aversion learning in rodents, where the onset of malaise minutes to many hours after tasting a substance leads to a robust and long-lasting taste avoidance memory [2]. More subtle cases involve learning with a CS-US interval of minutes or seconds [1,3]. Importantly, such learning requires that following cessation of the CS, a representation or 'trace' remains in the nervous system that can be functionally associated with a later US so that a memory is formed. Understanding how and where stimulus 'traces' manifest in the nervous system and how they are incorporated into memory is of

considerable interest. Animals with relatively simple and manipulable nervous systems may be useful for this endeavour.

Two recent studies [4,5] investigated trace conditioning using olfactory associative learning and physiological approaches in the honeybee and fruit fly. Bees quickly learn to extend their proboscis to odors following association of that odor with a sucrose reward [6,7]. In a similar manner, flies learn to avoid an odor that was previously associated with an electric shock [8]. Using variants of these established training protocols, both groups investigated parameters of 'trace' conditioning. Flies could learn a ten second odor stimulus when punished fifteen seconds afterwards, whereas bees could remember a half second odor pulse when rewarded six seconds later. Surprisingly, unlike other animals [3,4,9,10], bees remembered after only one trial of trace





Odor exposure activates unique combinations of olfactory sensory neurons (green). These neurons activate specific glomeruli in the antennal lobe engaging inhibitory (red) and excitatory (blue) local neurons and combinations of projection neurons (green). Projection neurons carry odor information to the mushroom body (MB) calyx where signals are transformed into activation of a small fraction (green) of the population of mushroom body neurons. Trace conditioning could utilize a lasting odor trace in mushroom body neurons (green). Adapted with permission from [20].

conditioning [5]. The ability to condition bees with a single trial indicates that learned associations likely involve a lingering CS 'trace' rather than a prediction error-driven adjustment of the reinforcement system [11].

One of the most interesting and apparently conserved features of trace conditioning is that performance can be improved by prior experience [9,10,12]. Successful trace conditioning with a particular time interval can extend the interval that can be bridged in subsequent trials. Conditioning bees or flies with a shorter interval between CS and US improved future learning with a time gap that was too long for naïve animals to learn. Interestingly, a previous trial of delay conditioning did not facilitate future trace conditioning in bees. It therefore appears that flies and bees benefit from being familiar with the

relevance of a gap between the stimuli. In effect they have recognized and learned one of the important 'rules' of the learning task so that they are able to maintain the utility of the CS trace for longer. Such a scenario is reminiscent of the concept of a preestablished mental schema or framework, which facilitate encoding, storage and consolidation of new associations in rats [13]. Establishing the neural mechanisms accounting for these, at least superficially, similar phenomena will be of great interest.

What are the neural correlates of the 'trace' that allow it to be associated with a temporally delayed US? Given that both the fly and bee learning tasks involve an olfactory CS, it seems likely that remnants of the odor representation will reside in the circuitry of the olfactory system (Figure 1). Indeed, prior work has indicated that odor responses remain in the bee or fly antennal lobe — the first center of insect olfactory processing — following the odor stimulus [14,15]. Furthermore, the similarity of odor responses in the antennal lobe is well correlated with the perceived similarity of odors following delay conditioning [16]. Both of the new studies [4,5] therefore used calcium (Ca²⁺) imaging to look for a CS trace in the bee or fly antennal lobe after the odor delivery was terminated.

Although the post-odor responses were reproducible and odor-specific. the similarity of their combinatorial profiles did not correlate with the odors perceived as similar by the animals following trace conditioning. Therefore, the authors [4,5] concluded that neither the primary olfactory receptor neurons of the fly nor the second order projection neurons of the bee maintained informative odor representations following the stimulus. Is this lack of a correlation enough to write off the neurons in the antennal lobe as potentially representing a CS trace? Certainly the data suggest that these Ca²⁺-dependent signals in the antennal lobe are not a good candidate to inform us of the nature of persistent odor traces in the nervous system. However, the antennal lobe circuitry might hold odor traces in a more complex temporal manner than that evident using Ca2+ imaging. Additionally, the odor traces might be represented amongst a larger antennal lobe neural ensemble and/or in Ca²⁺-independent intracellular signalling cascades within the relevant neurons. Lastly, given the apparent importance of the mushroom body for classical olfactory learning in the bee and fly, one might imagine the mushroom body to be the most likely site to locate prolonged associable odor traces that would also correlate to perceptual similarity. However, potential odor traces in the mushroom body might also be more complex than activity-evoked fluctuations of Ca²⁺ [17].

It is also conceivable that trace conditioning in bees and flies will involve neural circuitry that is not employed during routine delay conditioning. In mammals, trace conditioning recruits additional neural structures such as the hippocampus [10,18]. Although the reason is not fully understood, a simple explanation could be that the hippocampus is required to solve a more difficult task [3]. The hippocampal circuitry has been proposed to support a neuronal reverberation that holds the CS representation long enough for the animal to "bridge the gap" between the CS and US presentation. A conceptually similar mnemonic reverberant circuitry has been proposed to exist amongst subsets of fruit fly mushroom body neurons maintained by broadly arborizing excitatory dorsal paired medial neurons and inhibitory anterior paired lateral neurons [19]. It would be straightforward to test whether the function of this mushroom body circuitry is required to hold the odor trace and thereby permit olfactory trace conditioning in the fly. With this sort of analysis we might catch a glimpse of the elusive odor trace.

References

- Pavlov, I.V. (1928). Lectures on Conditioned Reflexes: The Higher Nervous Activity of Animals (Lawrence & Wishart, London: W.H. Gantt, Trans).
- Davis, C.M., and Riley, A.L. (2010). Conditioned taste aversion learning: implications for animal models of drug abuse. Ann. N. Y. Acad. Sci. 1187, 247–275.

- Beylin, A.V., Gandhi, C.C., Wood, G.E., Talk, A.C., Matzel, L.D., and Shors, T.J. (2001). The role of the hippocampus in trace conditioning: temporal discontinuity or task difficulty? Neurobiol. Learn. Mem. 76, 447–461.
- Galili, D.S., Ludke, A., Galizia, C.G., Szyszka, P., and Tanimoto, H. (2011). Olfactory trace conditioning in Drosophila. J. Neurosci. 31, 7240–7248.
- Szyszka, P., Demmler, C., Oemisch, M., Sommer, L., Biergans, S., Birnbach, B., Silbering, A.F., and Galizia, C.G. (2011). Mind the gap: olfactory trace conditioning in honeybees. J. Neurosci. 31, 7229–7239.
- Kuwabara, M. (1957). Bildung des bedingten Reflexes von Pavlovs Typus bei der Honigbiene, Apis mellifica. J. Fac. Sci. Hokkaido Univ. Ser. VI Zool. 13, 458–464.
- Bitterman, M.E., Menzel, R., Fietz, A., and Schafer, S. (1983). Classical conditioning of proboscis extension in honeybees (Apis mellifera). J. Comp. Psychol. 97, 107–119.
- Tully, T., and Quinn, W.G. (1985). Classical conditioning and retention in normal and mutant Drosophila melanogaster. J. Comp. Physiol. [A] 157, 263–277.
- Clark, R.E., and Squire, L.R. (1998). Classical conditioning and brain systems: The role of awareness. Science 280, 77.
- 10. Woodruff-Pak, D.S., and Disterhoft, J.F. (2008). Where is the trace in trace conditioning? Trends Neurosci. 31, 105–112.
- Schultz, W., Dayan, P., and Montague, P.R. (1997). A neural substrate of prediction and reward. Science 275, 1593–1599.
- Lucas, G.A., Deich, J.D., and Wasserman, E.A. (1981). Trace autoshaping: Acquisition, maintenance, and path dependence at long trace intervals. J. Exo. Anal. Behav. 36, 61–74.
- Tse, D., Langston, R.F., Kakeyama, M., Bethus, I., Spooner, P.A., Wood, E.R., Witter, M.P., and Morris, R.G. (2007). Schemas and memory consolidation. Science 316, 76–82.

- Sachse, S., and Galizia, C.G. (2002). Role of inhibition for temporal and spatial odor representation in olfactory output neurons: a calcium imaging study. J. Neurophysiol. 87, 1106–1117.
- Silbering, A.F., Okada, R., Ito, K., and Galizia, C.G. (2008). Olfactory information processing in the Drosophila antennal lobe: anything goes? J. Neurosci. 28, 13075–13087.
- Guerrieri, F., Schubert, M., Sandoz, J.C., and Giurfa, M. (2005). Perceptual and neural olfactory similarity in honeybees. PLoS Biol. 3, e60.
- Ito, I., Ong, R.C., Raman, B., and Stopfer, M. (2008). Sparse odor representation and olfactory learning. Nat. Neurosci. 11, 1177–1184.
- Solomon, P.R., Vander Schaaf, E.R., Thompson, R.F., and Weisz, D.J. (1986). Hippocampus and trace conditioning of the rabbit's classically conditioned nictitating membrane response. Behav. Neurosci. 100, 729–744.
- Pitman, J.L., Huetteroth, W., Burke, C.J., Krashes, M.J., Lai, S.L., Lee, T., and Waddell, S. (2011). A pair of inhibitory neurons are required to sustain labile memory in the Drosophila mushroom body. Curr. Biol. 21, 855–861.
- Keene, A.C., and Waddell, S. (2007). Drosophila olfactory memory: single genes to complex neural circuits. Nat. Rev. Neurosci. 8, 341–354.

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Adaptive Introgression: The Seeds of Resistance

Populations of the European house mouse have acquired resistance to anticoagulant pesticides from a closely related species. This discovery improves our understanding of the circumstances in which interspecific genetic exchange is likely to facilitate adaptation.

Loren Rieseberg

My favorite paper as a graduate student was a review by botanist, Charley Heiser [1], titled "*Introgression re-examined*", in which he evaluated the role of interspecific gene exchange (introgression) in evolution. Botanists had long viewed introgression as a potent evolutionary force that promoted the development and acquisition of novel adaptations [2,3]. However, zoologists were more skeptical and typically emphasized the frequently negative fitness consequences of hybridization [4]. Despite his botanical background, Heiser [1] was forced to conclude that while introgression "may play a very significant role; it must be admitted, there is as yet no strong evidence to support such a claim."

The strong evidence Heiser was looking for has been a long time coming. Molecular marker studies over the past three decades have shown that introgression is widespread in plants and animals [5–7], vindicating earlier botanical views about the porous nature of reproductive barriers [1–3,8]. However, convincing evidence of adaptive introgression has been more difficult to obtain. The paucity of examples might be due to the challenge of marshalling the diverse data sets required to demonstrate that introgressed alleles or traits have been favored by selection [9–11]. Alternatively, it might be that adaptive introgression is rare. A paper by Michael Kohn and colleagues [12] in this issue of *Current Biology* not only furnishes exceptionally strong evidence of adaptive introgression involving Old World mice, but it also provides insights into the circumstances under which introgression is likely to facilitate adaptive evolution.

Theory indicates that for many species — especially those with large populations — variation created by mutation is likely to exceed that provided by introgression [13]. Thus, introgression is thought to promote adaptation mainly through the transfer of favorable sets of genes rather than individual mutations. A familiar example of the latter — albeit involving horizontal gene transfer rather than sexual hybridization and