

For example, could the auditory cortex recover after the 'retardation' and eventually function as it does in normal adults? What are the cellular and molecular bases that would allow the putative reopening or prolongation of the critical period? The end of the critical period in cortical development of normal animals has been shown to correlate with the changes in certain neurotransmitter receptor properties (e.g. NMDA receptors) as well as with the level of brain-derived neurotrophic factor (BDNF) [19–21]. It would be interesting to know whether behavioral manipulations used by Chang and Merzenich altered underlying molecular signaling mechanisms in auditory cortex. Studies of these questions will help us understand whether the extended cortical plasticity observed in noise-reared animals is an extension of the critical period that is observed in normally raised animals or whether it constitutes another form of cortical plasticity.

Nevertheless, the notion of a 'plastic critical period' has tremendous implications. Studies along this line could uncover unified mechanisms that link all types of cortical plasticity over the lifespan of an animal or human. Such findings should also have therapeutic implications for treating children with hearing loss at young ages. If the outcome of the Chang and Merzenich study and of studies of electrical stimulation in congenitally deaf cats [11] are of any indication, methods to help regain normal function of auditory cortex, and consequently speech and language processing ability, in affected children are some of the promising possibilities just over the horizon.

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## Dissecting complex behaviours in the post-genomic era

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Identifying gene functions in behaviours has so far relied mainly on achievements in the field of molecular genetics. Further progress can be made by developing

new approaches that allow refinement of behavioural phenotypes. The current availability of several thousand different mutant mice challenges behavioural neuroscientists to extend their views and methodologies, to dissect complex behaviours into behavioural phenotypes, and subsequently to define gene-behavioural

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### phenotype relationships. Here, we plead for multi-day automated behavioural observations in carefully designed environments.

Behavioural testing in animals, a crucial feature of phenotyping in neuroscience, is usually based on measuring behavioural responses to environmental events that are induced by the experimenter. The frequently used open-field test, for example, is commonly employed to study general activity and fear-related behaviours in mice and rats [1,2]. In this test, movements of the animal are monitored up to one hour after the animal has been placed in a novel open arena from which it cannot escape. Additional tests, such as the elevated plus maze [3] and the light–dark box [4], allow external validity of the observed open-field behaviours. However, these tests are short-lasting, and depend on individual locomotor activity levels and novelty responsiveness of the animal, as well as on human interference. This hampers their use for determining gene–behavioural phenotype relationships and stresses the need for new analytical procedures addressing the complex behaviours. Although some ideas for overcoming these problems have been put forward, such as improving currently available tests, using test batteries and increasing test information density [5,6], behavioural complexity and gene–environment interactions require new methodology in this field of research.

### Interacting physiological processes

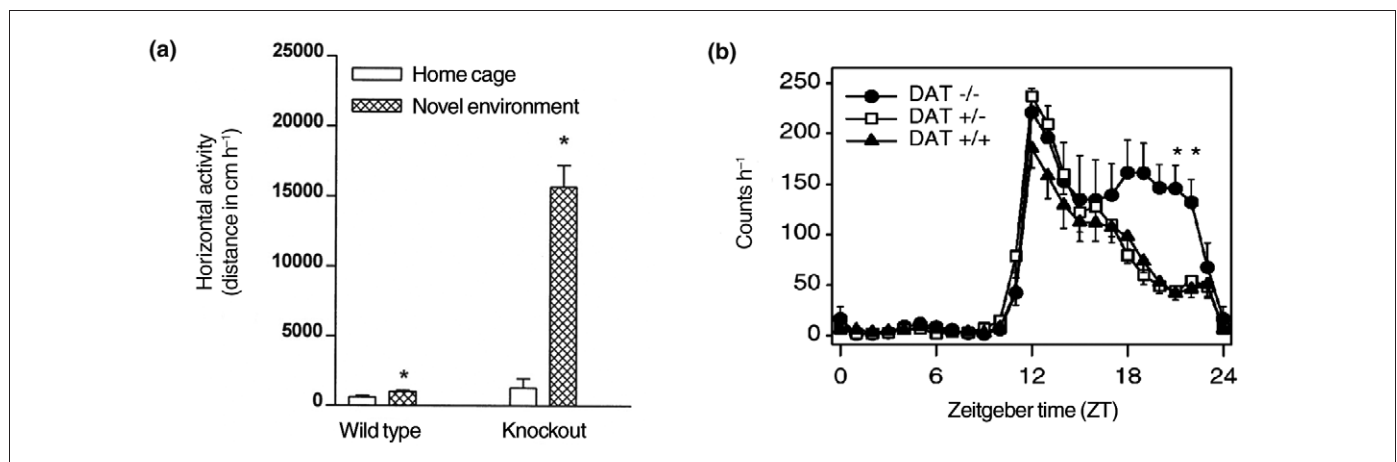
Expression of a particular behaviour at a certain time is triggered by the integration of internal and external signals (such as hunger and food availability, respectively) and is guided by the ability of the animal to execute proper behavioural responses. For instance, a hungry mouse that searches for new food resources relies on an efficient exploration strategy in which finding the food resource in time and taking the risk of being exposed to predators need

to be balanced. Furthermore, in the wild, mice face the risk of using more energy to gather food than the obtained food gives them in return on any given day. Because exploration for food is influenced by different integrated physiological processes (e.g. energy balance, motor action and fear), as well as by environmental factors (e.g. variations in ambient temperature, in food availability and in photoperiod), the design of behavioural laboratory methods that dissociate the various behavioural components is a challenge. Although limitations in behavioural testing environments and duration are practical realities of the mouse laboratory, these parameters should be carefully chosen and directed by a clearly defined working hypothesis.

Conventional laboratory tests, such as the open-field arena, touch upon different aspects of exploratory behaviour, such as locomotor activity and fear-related processes. However, during the relative short testing episode generally employed, it is impossible to discriminate between gene function in novelty-induced and baseline behaviours. For example, mice that lack the dopamine transporter gene have behavioural locomotor activity levels under baseline conditions that are comparable to those in wild-type animals, but they exhibit a >12-fold increase of locomotion following placement in a novel environment. Although dissociation of novelty-induced and baseline locomotor behaviour in this mutant was observed during a relative short-lasting testing procedure, the time of day that these tests are performed can highly influence the outcome of the observations [7–9] (Figure 1). Thus, characterization of gene functions in exploratory behaviour requires behavioural paradigms that allow dissection of this complex behaviour into different components in view of circadian-induced variations of these components.

### Interference and order effects

Executing multiple behavioural tests usually involves experimenter interference, such as handling or transport



**Figure 1.** Dissociation of gene function in behavioural regulation requires assessment of novelty-induced and baseline behaviours during the appropriate circadian phase. Although brief assessments of behaviour (e.g. for 1 h) are attractive with respect to the screening capacity, they can yield discrepancies with longer assessments of behavioural testing (e.g. 24 h). For example, hyperactivity in mice that lack the gene encoding the plasma membrane dopamine transporter (DAT) appeared to be novelty driven because locomotor activity was ~12-fold higher in an environment that was novel to the mouse at the start of the experiment (a). Asterisks indicates  $P < 0.05$  for activity in familiar versus novel environments. No significant differences between the activities of DAT-knockout and wild-type mice were observed in their home cages during the 1 h of behavioural monitoring. (b) In another study, 24-h rhythms in behavioural activity were measured in DAT  $-/-$ , DAT  $+/-$  and wild-type (DAT  $+/+$ ) mice. Locomotor activity levels were higher in DAT  $-/-$  animals, but only during the last part of the dark phase while housed in home cages. Thus, continuously behavioural observations can be crucial for gene function characterization in novelty-induced and baseline behaviours that can vary across different phases of the 24-h cycle. Zeitgeber time (ZT) 0–12 and ZT 12–24 refer to the 12-h light and the 12-h dark phases of the 24-h light–dark cycle, respectively. Panel (a) reproduced, with permission, from Ref. [8] © (1999) American Association for the Advancement of Science; (b) reproduced, with permission, from Ref. [7] © 2001 by the Society for Neuroscience.

of animals [10], and cues from the experimenter that influences the behavioural performance of an animal [11] (Figure 2). For example, measuring pain responses in mice revealed that experimenter effects account for more trait variability than genotype [12]. In addition, recent studies have shown that simultaneous testing of several inbred mice strains and a mutant mouse strain across three laboratories resulted in behavioural expression differences, despite standardization of the test procedures [13]. Mice exposed to a battery of various behavioural tests expressed significant lower levels of locomotor activity in the open field than mice that were naïve to behavioural testing [14]. These order effects could even be amplified in animals with selective mutations in genes that are involved in physiological processes, such as coping strategies to changing environments. Circumvention of these interfering procedural aspects is required to reduce nonspecific environmental influences on the gene–behavioural phenotype relationship.

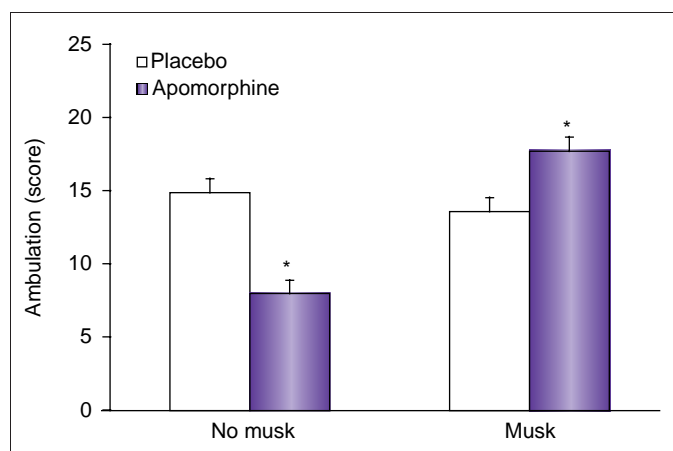
### Dissection of behavioural phenotypes

Studies in the field of biological rhythms have revealed that behavioural observations during several consecutive days or weeks in the home cage of an animal allow reliable assessment of stable behavioural circadian rhythms that are highly sensitive to environmental signals, such as light and human interference [15,16]. Because behavioural observations during several days can also dissociate novelty-induced and baseline behaviours at different phases of the light–dark cycle, behavioural monitoring in the home cage will significantly contribute to the refinement of behavioural phenotypes. In addition, by carefully designing a home cage environment that addresses different behavioural characteristics of interest, complex behaviours can be further dissected into behavioural phenotypes with minimum human interference. For example, place preference and the expression of locomotor activity can be simultaneously measured by introducing shelter places in the home cage. Other behavioural parameters, such as

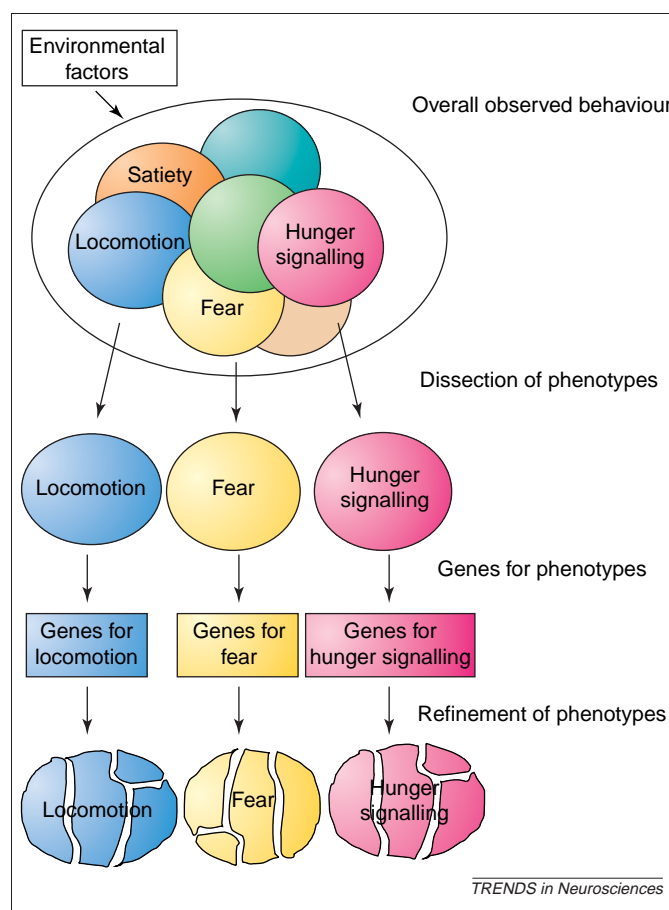
eating behaviour, can also simultaneously be dissected by recording meal frequency and duration. One should keep in mind, however, that some environmental factors can mask the expression of particular behaviours. For example, access to running wheels is often introduced as an easily detectable measure for circadian rhythms and locomotor activity levels in rodents, but wheel-running activity can significantly alter the amplitude of behavioural rhythms [17]. Thus, behavioural monitoring, including automated scoring, for multiple days in a home cage environment without confounding factors (e.g. masking effects), will further contribute to the refinement of behaviour and dissection of complex behaviours into different behavioural phenotypes within pre-defined working hypotheses.

### Concluding remarks

In the post-genomic era, studies about the relationship between genotype and behavioural phenotype require novel methodology to measure behaviours. Because



**Figure 2.** Experimenter interference during behavioural assessment is a major confounding factor on the outcome of a behavioural test [12] and can even reverse the phenotype. Injection of a low dose of the dopamine agonist apomorphine (10 ng) into the nucleus accumbens of the brain decreased locomotor activity in rats (left). When the experimenter applied musk perfume to their wrists on the evening before the experiment, apomorphine induced hyperactivity (right) instead of hypoactivity. Asterisks indicate a significant different from placebo treatment. Reproduced, with permission, from Ref. [11].



**Figure 3.** Determination of gene–behavioural phenotype relationships first requires dissection of the overall observed behaviour into behavioural phenotypes. The expression of the overall observed behaviour is likely to result from a complex gene–environment interaction [e.g. of molecules that regulate hunger signalling, fear, locomotion and other behavioural components (coloured ovals) with environmental factors such as food availability, ambient temperature and photoperiod variation]. Development of paradigms that allow dissection of complex behaviours into different behavioural phenotypes will optimize the search for genetic determinants for these phenotypes. Because genes influence brain region-specific aspects of a certain behavioural phenotype [20,21], local gene expression or deletion technology can further refine phenotypes when studied in environments closely related to the natural territory of the animal. In this way, a dissected phenotype such as hyperactivity can, for example, be further refined with respect to hyperactivity in nest-building behaviour or in food exploratory behaviour [21].

behavioural expression relies heavily on many interactions between internal and external factors, working hypotheses should direct the balance between environmental conditions and recording duration, taking into account the practical feasibility within the laboratory and the behavioural resolution that minimizes false-positive and false-negative findings. Inspired by the successful identification of novel genes involved in circadian rhythms [18,19], the use of automated behavioural measurements for several consecutive days in the home cage of an animal is put forward as a way to reduce significant contributions of laboratory-specific variation and experimenter interference. This approach would also contribute to the dissociation of interacting physiological processes, such as novelty-induced and baseline behavioural activity, as well as circadian variation in behavioural expression. Multi-day recordings will, as a matter of course, reduce the behavioural screening rate of a mouse colony. But, with the improvements of behavioural resolution, it is also worth considering a home cage environment that assesses all behavioural components of interest at once, in contrast to a behavioural screening program that entails multiple short-lasting tests each addressing a single behavioural component. Whether the focus of interest is cognition, social activity, anxiety or eating, the development of well-designed home-cage environments offers great opportunities for neuroscientists to optimize the dissection of complex behaviours into behavioural phenotypes, and the subsequent identification of the genes for these phenotypes (Figure 3). Complex behaviour is influenced by multiple genes, and one gene generally affects several aspects of a complex behaviour. Interestingly, recent studies on brain region-specific gene expression have shown that a single gene can exert multiple functions within a behavioural domain depending on the brain region [20,21]. Thus, rodents with local vector-directed gene expression or conditional knockout mouse lines monitored in environments that address additional aspects of their natural habitat (e.g. temperature and photoperiod variations) will further optimize the search for gene-behavioural phenotype relationships.

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