

Molecular Separation of Two Behavioral Phenotypes by a Mutation Affecting the Promoters of a Ca-Activated K Channel

Nigel S. Atkinson, Robert Brenner, Whei-meih Chang, Jennette Wilbur, James L. Larimer, and Joyce Yu

Section of Neurobiology, School of Biological Sciences and the Institute for Cellular and Molecular Biology, The University of Texas at Austin, Austin, Texas 78712-1064

The *Drosophila* *slowpoke* gene encodes a BK-type calcium-activated potassium channel. Null mutations in *slowpoke* perturb the signaling properties of neurons and muscles and cause behavioral defects. The animals fly very poorly compared with wild-type strains and, after exposure to a bright but cool light or a heat pulse, exhibit a “sticky-feet” phenotype. Expression of *slowpoke* arises from five transcriptional promoters that express the gene in neural, muscle, and epithelial tissues. A chromosomal deletion (*ash2*¹⁸) has been identified that removes the neuronal promoters but not the muscle–tracheal cell

promoter. This deletion complements the flight defect of *slowpoke* null mutants but not the sticky-feet phenotype. Electrophysiological assays confirm that the *ash2*¹⁸ chromosome restores normal electrical properties to the flight muscle. This suggests that the flight defect arises from a lack of *slowpoke* expression in muscle, whereas the sticky-feet phenotype arises from a lack of expression in nervous tissue.

Key words: *Drosophila*; calcium-activated potassium channel; potassium channel; behavior; flight; tissue-specific transcription; regulation of transcription

Ion channel proteins generate the electrical impulses used by neurons and muscles to convey information and trigger movement. The range of electrical properties that a cell can manifest arises from the combined activity of the suite of channels expressed. Ion channels also participate in processes distinct from the transmission of electrical signals. In many epithelial cells, the same family of channels participate in transport of ions, water, and nutrients between the lumen of organs and their interior (Pacha et al., 1991; Stoner and Morley, 1995). The combination of channels expressed is expected to be substantially different in these functionally disparate cells.

The choice of which channels a cell is to express does not appear to be a simple decision. The superfamily of ion channels are represented by a large number of distinct genes, some of which can encode multiple products via alternative promoter use and alternative splicing (Brenner and Atkinson, 1996; Wei et al., 1996). In general, the number of biophysically distinct channels expressed by a cell is small compared with its potential.

The *Drosophila* *slowpoke* gene encodes a Ca-activated K channel expressed in neurons, muscles, midgut, and trachea (Becker et al., 1995). Elimination of the channel by mutation dramatically alters the electrical properties of both neurons and muscles (Elkins and Ganetzky, 1988; Saito and Wu, 1991; Warbington et al., 1996). In tracheal and midgut cells, the function of the channel has not been directly demonstrated, but it is believed to participate in the process of electrolyte transport and acid secretion, respectively (Becker et al., 1995; Brenner and Atkinson, 1997). It is unlikely that a single channel polypeptide satisfies the functional needs of such disparate tissue types. The *slowpoke* gene, however, is well suited for this role. Expression of *slowpoke* arises

from an array of tissue-specific transcriptional promoters, some of which give rise to mRNAs that encode polypeptides differing in their N terminus. To date, five tissue-specific promoters have been mapped (Bohm et al., 2000). In addition, *slowpoke* transcripts are alternatively spliced at five sites that affect the coding region of the gene (Atkinson et al., 1991; Adelman et al., 1992). It is assumed that multiple promoters and alternative splicing enable tissues to express channels tailored to the needs of the cell.

In addition to their electrophysiological phenotypes, *slowpoke* mutants display behavioral abnormalities. The animals are semi-flightless and, in response to a brief heat pulse, remain stationary for many minutes (Elkins et al., 1986). This unusual temperature-dependent phenotype is difficult to understand in terms of thermolabile proteins. Here, we demonstrate that the animals are not paralyzed but inappropriately adhere to the surface and that this behavior is not wrought by temperature per se but appears to be a consequence of overstimulation. Using a chromosomal deletion that removes the neuronal but not the muscle promoter, we demonstrate the origins of the “sticky-feet” and the flight phenotypes.

MATERIALS AND METHODS

Stocks. *Drosophila* stocks were maintained using standard *Drosophila* husbandry techniques. Four stocks were used: *w*¹¹¹⁸; *st hh*^{bar3} *slo*¹, *slo*⁴, *y w*; *red e ash2*¹⁸/*y+* TM3 *Sb e Ser*, and *red, e, ash2*¹⁸/TM6 *Tb*. The *ash2*¹⁸ allele stock was kindly provided by Allen Shearn (Department of Biology, The Johns Hopkins University, Baltimore, MD).

Reverse transcription-PCR. For reverse-transcription (RT)-PCR, RNA was purified from *w*¹¹¹⁸ flies, from *slo*⁴ flies, and from *red e ash2*¹⁸/*slo*⁴ transheterozygotes. Approximately 0.5 gm of adult animals were added to 2.5 ml (5 vol) of 3 M LiCl, 6 M urea, and 0.2% SDS and ground in a glass homogenizer. To precipitate the RNA, the sample was incubated overnight on ice and centrifuged at 5000 rpm [Sorvall (Newtown, CT) RC5C with a SA-600 rotor] for 20 min at 4°C in a 15 ml Corex centrifuge tube. The liquid between the bottom pellet and the floating pellet was removed with a pipette and discarded. The pellet was dissolved in 2–4 vol of 10 mM Tris-base, 1 mM Na₂EDTA, and 1% SDS on ice for 15 min. The solution was extracted twice with phenol:CCl₃ (24:1), pH 8.0 with 0.1% hydroxyquinoline, and once with CCl₃. The RNA was precipitated by adjusting the aqueous layer to 0.3 M sodium acetate (stock is pH 5.2),

Received Dec. 22, 1999; revised Dec. 22, 1999; accepted Feb. 2, 2000.

This work was supported by National Science Foundation Grant IBN-9724088 to N.S.A.

Correspondence should be addressed to Nigel S. Atkinson, Section of Neurobiology, Patterson Building, The University of Texas at Austin, Austin, TX 78712-1064. E-mail: nigela@mail.utexas.edu.

Copyright © 2000 Society for Neuroscience 0270-6474/00/202988-06\$15.00/0

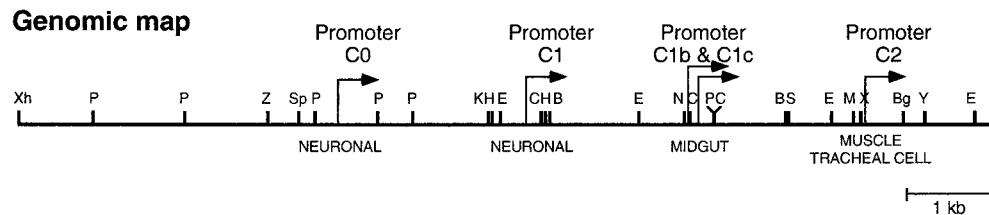


Figure 1. Map of the *slowpoke* transcriptional control region. The rightward-pointing arrows identify the position of five *slowpoke* transcriptional promoters. The labels immediately below the line identify the tissue specificity of the each promoter as determined by deletion mapping. A, *Apa*I; B, *Bam*HI; Bg, *Bgl*II; C, *Cl*aI; E, *Eco*RI; H, *Hind*III; K, *Kpn*I; M, *Mun*I; N, *Nco*I; P, *Pst*I; S, *Sma*I; Sp, *Spe*I; X, *Xba*I; Xh, *Xho*I; Y, *Xmn*I; Z, *Sph*I.

followed by the addition of 3 vol of 100% ethanol and incubation at -70°C . The RNA was stored in ethanol until needed. To recover the RNA, the solution was centrifuged at 8500 rpm/30 min/ 4°C . The pellet was air-dried and resuspended in 300–600 μl of water. All solutions were made using DEPC-treated water.

One microgram of RNA was reverse-transcribed in a 25 μl volume with Moloney Murine Leukemia Virus Reverse transcriptase (New England Biolabs, Beverly, MA) and 0.5 μM of the primer slo26 (5' TGCGATC-CAGTATGCAGTCT 3') for 45 min at 40°C . The solution was adjusted to 100 μl with water and heated at 65°C for 5 min. Five microliters of the RT product was used to seed the PCR reaction.

A hot-start PCR amplification was performed in a PCR machine (MJ Research Inc., Watertown, MA) with the enzyme mix provided in a 100 μl volume using the enzyme mix provided in the Expand High Fidelity PCR System (Boehringer Mannheim, Indianapolis, IN) and 1.5 mM MgCl₂, 0.125 mM dNTPs, and 2 μM of each PCR primer. Annealing temperatures were determined using the OLIGO program (National Biosciences, Plymouth, MA).

To detect transcripts produced by *slowpoke* promoter C0, PCR was performed using the GAMMA5 primer (5' ATTGTATACGCTGCT-GACGAGA3', anneals to exon C0) and slo45 primer (5' CCGCCATTT-TGATTCTGTGTG3', anneals at approximately nucleotide 2520). The slo42 primer (5' CTCGGTGGTTTAGCCAGTACTA 3'), which anneals to exon C1, and the slo45 PCR primer were used to detect the presence of transcripts produced from both promoter C0 or promoter C1. Neither of these primers amplify a product derived from transcripts produced from promoter C2. Promoter C2 products were detected by using the PCR primer slo43b (5'TGGCACTCGACTGCACTTGA3') and primer slo45. The slo43b primer specifically anneals to exon C2, and therefore this primer set can be used to detect transcripts produced by this promoter C2.

Action potential recording. The recording of action potentials from the dorsal longitudinal flight muscles (DLM) was performed essentially as described by Elkins et al. (1988). Lightly etherized adults were glued to a coverslip along their ventral midline using Superglue. Flies were allowed to recover for a minimum of 1 hr before recording. The mounted fly was placed under a microscope, and two uninsulated tungsten electrodes were inserted into the brain. Current flow [generated with a Grass Instruments (Quincy, MA) S88 stimulator] through these electrodes causes activation of the giant fiber pathway and the production of action potentials in the DLM. A glass electrode was inserted into the flight muscle (2–5 M Ω , filled with 1 M NaCl) and used to record the evoked action potential spikes using a World Precision Instruments (Sarasota, FL) Electrometer Intra 767. All recordings were made in DLM indirect flight muscles c through f (Engel and Wu, 1992). All potentials were measured with reference to an uninsulated tungsten electrode inserted into the abdomen. The stimulus threshold was determined by stimulating the brain at 2 V (0.1 msec) and gradually increasing the potential until DLM spikes were observed. After a threshold voltage was determined, 0.2 V above threshold was used for the remainder of the experiment. Data were collected using a MacADIOS 8ain analog-to-digital converter (69.4 μsec /point), filtered at 14 kHz with a single-pole low-pass filter, and recorded using the Macintosh program Superscope (GW Instruments, Somerville, MA).

Sticky-feet behavioral assay. The sticky-feet behavior is elicited by overstimulating the flies using a heat pulse or a bright light delivered by a fiber optic lamp. For heat treatment, 2- to 5-d-old adult flies are trapped at the bottom of an empty glass fly vial using a foam or cotton plug and incubated at $37.7\text{--}40^{\circ}\text{C}$ for 2–8 min. The time and temperature required to elicit the behavior seems to vary with the season but not within a

season. Exposure for a few seconds to a very bright but cool light can be substituted for the heat treatment (50 W fiber optic lamp, set on high). Positive (wild-type animals) and negative (*slo*⁴ mutant animals) internal controls are always performed. After the heat pulse, the animals are gently transferred to the tabletop and not disturbed for ~ 15 sec. A pencil with a pink-pearl eraser was used to push on the sides of the animals. Flies homozygous for a *slowpoke* mutant allele hang onto the surface and allow themselves to be pushed over. Flies heterozygous for a *slowpoke* mutant allele will walk or fly away from the stimulus. Wild-type flies will take flight and leave the area.

Flight test. The relative ability of the 3- to 6-d-old animals to fly was measured as described previously (Benzer, 1973; Elkins et al., 1986; Green et al., 1986) with minor modifications. The walls of a pipette jar (15 cm diameter, 62 cm tall) were coated with mineral oil. A funnel was fixed on a platform at the top of the jar, and flies were dropped through the opening. The falling animals fly toward the walls of the jar and are trapped in the mineral oil. The position of each animal on the jar is marked, and then the distance of each mark from the bottom of the jar is determined. Rulers taped to the side of the jar simplify this process. Although the mineral oil seeps to the bottom of the jar, the embedded flies do not move and are stable for hours. Animals that fly well tend to cluster near the top of the jar, whereas animals that fly poorly are most often found near the bottom. Dead flies and flies that do not fly fall into the bottom of the jar and are not counted. This test can be used to distinguish wild-type animals from animals carrying a mutant *slowpoke* allele (Elkins et al., 1986). For each genotype, 350–1000 flies were tested. For flight testing, animals were not heat-treated.

RESULTS

Fine mapping of *ash2*¹⁸ deletion endpoint on the *slowpoke* promoter map

The *slowpoke* transcriptional control region has been well characterized using transgenes that drive the expression of a reporter gene. It has been shown that promoter C0 and C1 (Fig. 1) generate *slowpoke* expression in the nervous system in adult, larval, and embryonic stages, whereas promoter C2 alone is responsible for expression in muscle and tracheal cells (Bohm et al., 2000).

The *slowpoke* gene is found on chromosome 3 at cytological position 96A17 (Atkinson et al., 1991). Adjacent to *slowpoke* is the *ash2* gene in which mutations cause homeotic transformations of body parts during development. The function of the *ash2* gene and the phenotype of its mutant alleles are unrelated to that of the *slowpoke* gene. However, during the study of *ash2* function, Adamson and Shearn (1996) characterized a chromosomal deletion that removed the entire *ash2* gene. This mutation is called *ash2*¹⁸. Fortunately, one endpoint of this encroached upon the *slowpoke* gene. Their data indicated that this deletion removed some of the *slowpoke* transcriptional promoters.

Our mapping data indicates that the endpoint of the deletion falls between promoter C1 and promoter C2 (data not shown) and as a consequence removes promoter C0 and promoter C1 (Fig. 1). Deletion analysis performed on a *slowpoke* transgene has shown that removal of these two promoters causes a loss of

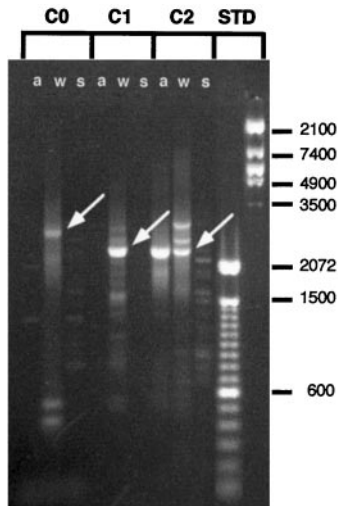


Figure 2. The *ash2*¹⁸ deletion eliminates expression from neuronal promoters C0 and C1 but not from the muscle–tracheal cell-specific promoter C2. The products of each promoter begins with a unique exon and can be identified by RT-PCR using exon-specific primers. Total RNA from *ash2*¹⁸/*slo*⁴ transheterozygous, wild-type, and *slo*⁴ homozygous flies was reverse transcribed, and the resultant cDNA was subjected to the PCR using the exon-specific primers. PCR products were separated in a 2% agarose gel and stained with ethidium bromide. C0, C1, and C2 identify groups of three lanes displaying products amplified using primers specific for exon C0, C1, or C2, respectively. Amplifications performed on *ash2*¹⁸/*slo*⁴, wild-type, and *slo*⁴ RNA are identified by *a*, *w*, and *s*, respectively. From left to right, the arrows identify PCR products diagnostic for the presence of mRNAs that include exon C0, exon C1, and exon C2, respectively. Amplification with exon C0 primers produces the 2844 nucleotide band only from wild-type RNA (lanes 1–3). The exon C1-specific primers produce the diagnostic 2406 nucleotide PCR product only from the wild-type RNA (lanes 4–6). Finally, the exon C2-specific primers amplified the diagnostic 2373 nucleotide product derived from exon C2 from both *ash2*¹⁸/*slo*⁴ and wild-type RNA but not from the *slo*⁴ RNA. Other bands are nonspecific PCR artifacts. Conditions favoring maximal sensitivity often lead to the concomitant production of spurious bands.

slowpoke expression in the adult CNS. Muscle and tracheal cell expression, however, persists if promoter C2 and the following downstream intron are present (Brenner and Atkinson, 1996; Brenner et al., 1996). Thus, the *ash2*¹⁸ deletion has removed sequences absolutely required for adult neuronal expression but left the promoter driving muscle and tracheal expression intact.

An RT-PCR assay was used to confirm that the *ash2*¹⁸ chromosome does not express the promoter C0 and C1 neuronal transcripts but does express transcripts originating from promoter C2. Each of these promoters produces a transcript that begins with a unique exon. RNA was isolated from *ash2*¹⁸/*slo*⁴ transheterozygotes. The use of the transheterozygote was necessary because the *ash2*¹⁸ deletion is a recessive lethal mutation. The *slo*⁴ mutant allele is a chromosome inversion with one breakpoint in the *slowpoke* gene. It has been shown to be a null mutation by genetic, electrophysiological, immunohistochemical, and RT-PCR criteria (Atkinson et al., 1991; Becker et al., 1995). RT-PCR was performed on wild-type, *ash2*¹⁸/*slo*⁴, and *slo*⁴ RNA using primer sets specific for transcripts containing exon C0, C1, and C2 (Fig. 2). Exon C0- and C1-containing transcripts were not detected in either the *ash2*¹⁸/*slo*⁴ or the *slo*⁴ RNA. A primer set specific for exon C2-containing transcripts amplified a product from *ash2*¹⁸/*slo*⁴ RNA but not from *slo*⁴ RNA. All of the primer sets amplified a product from the wild-type RNA.

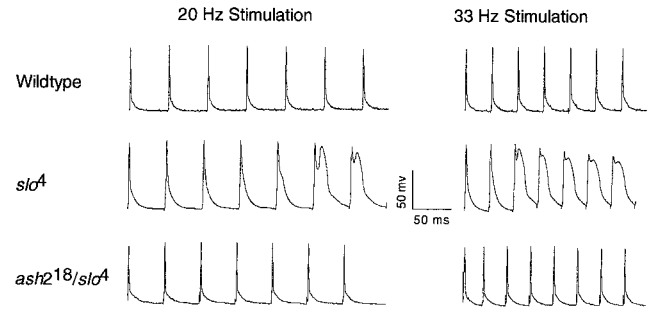


Figure 3. Action potentials produced by dorsal longitudinal flight muscles of wild-type, *slo*⁴ homozygous, and *ash2*¹⁸/*slo*⁴ transheterozygous animals at a stimulation frequency of 20 and 33 Hz. Wild-type muscle produces sharp action potentials at all stimulation frequencies tested. Muscle homozygous for the *slo*⁴ mutation initially produces sharp action potentials, but later spikes are of abnormal shape and breadth. The *ash2*¹⁸ deletion completely complements the *slo*⁴ action potential phenotype. This indicates that the *ash2*¹⁸ deletion produces functional *slowpoke* channels in *Drosophila* muscle.

Action potential recordings

In *Drosophila* muscle, the rising phase of the regenerative action potential is generated by the influx of Ca ions through voltage-gated Ca channels (Salkoff and Wyman, 1983). The repolarization phase of the action potential is driven by the activity of at least four different outward potassium currents. These are called I_A , I_{CF} , I_K , and I_{CS} . The *Shaker*, *slowpoke*, and *shab* genes encode the channels that conduct I_A , I_{CF} , and I_K , respectively (Baumann et al., 1987; Kamb et al., 1987; Papazian et al., 1987; Tsunoda and Salkoff, 1995; Singh and Singh, 1999). The channel that conducts I_{CS} has not yet been identified.

The electrophysiological consequences of a mutant *slowpoke* gene have been extensively studied. Mutant *slowpoke* alleles eliminate the Ca-activated K current called I_{CF} in neurons and muscle fibers (Salkoff, 1983; Elkins et al., 1986; Gho and Mallart, 1986; Singh and Wu, 1989; Komatsu et al., 1990; Saito and Wu, 1991; Brodie and Bate, 1993). In the indirect flight muscles of the adult, the loss of this potassium current has a striking electrophysiological consequence: the production of extremely broad and Bactrian camel-shaped action potentials (Elkins and Ganetzky, 1988). All of the mutant *slowpoke* alleles, *slo*¹, *slo*², *slo*³, *slo*⁴, *slo*⁵, and *slo*⁸, have been shown to eliminate I_{CF} in indirect flight muscles and to cause the same action potential phenotype (Atkinson et al., 1991).

To demonstrate that the *ash2*¹⁸ chromosome produced functional *slowpoke* channels in *Drosophila* flight muscle, we examined the shape of action potentials produced by the indirect flight muscles of *ash2*¹⁸/*slo*⁴ transheterozygotes. It was not possible to assay *ash2*¹⁸ in the homozygous state because it is a recessive lethal mutation. The *slo*⁴ mutation is a chromosomal inversion and has been shown to be a homozygous viable *slowpoke* null allele (Atkinson et al., 1991; Becker et al., 1995). Figure 3 shows action potentials produced from wild-type muscle, *slo*⁴ homozygous muscle, and *ash2*¹⁸/*slo*⁴ muscle stimulated at 20 and 33 Hz. In addition, we also examined action potentials evoked at 10 and 50 Hz (Fig. 4). In all cases, wild-type and *ash2*¹⁸/*slo*⁴ animals produced action potentials of normal shape and duration, whereas homozygous *slo*⁴ muscle produced broad and abnormally shaped action potentials. Therefore, in indirect flight muscle, the *ash2*¹⁸ chromosome produces functional *slowpoke* channels.

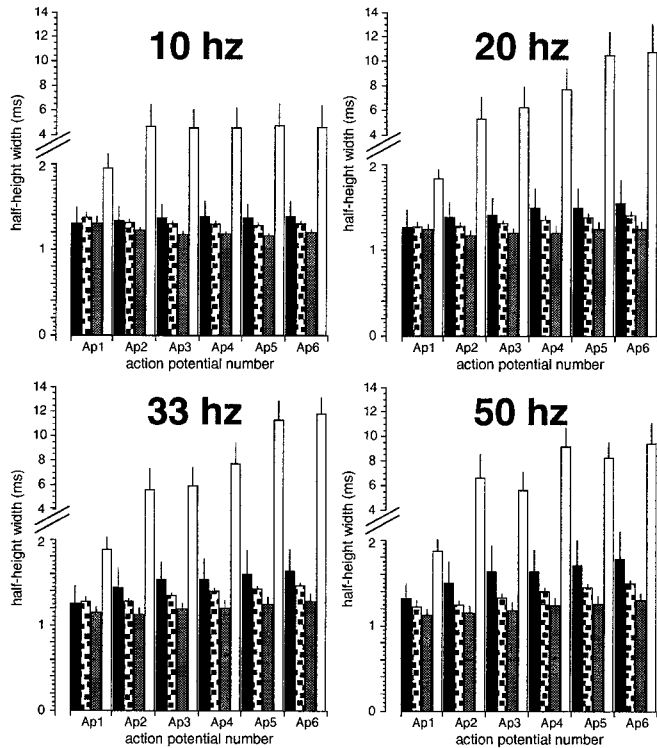


Figure 4. Broadening of dorsal longitudinal flight muscle action potentials stimulated at 10, 20, 33, and 50 Hz. The half-height width of six consecutive action potentials were measured. Width is expressed in milliseconds. *Black, stippled, gray, and white* bars represent an average from wild-type animals ($n = 3$), $slo^4/+$ heterozygote animals ($n = 9$), $ash2^{18}/slo^4$ transheterozygous animals ($n = 6$), and slo^4 homozygous animals ($n = 10$), respectively. The *thin line* above the bar represents the SEM for each measurement. *Ap1, Ap2, Ap3, Ap4, Ap5, and Ap6* represents the width of the first, second, third, fourth, fifth, and sixth action potentials, respectively.

Flight assay

A second phenotype caused by a defect in the *slowpoke* gene is a reduced ability of the mutant animals to fly (Elkins et al., 1986). Upon visual inspection, it is obvious that flies that carry a null mutation in the gene are very poor fliers. The *slowpoke* mutant animals walk much of the time and fly for only very short distances. We tested the capacity of the animals for flight using a modified cylinder drop test assay (Benzer, 1973; Elkins et al., 1986; Green et al., 1986). In this assay, flies are dropped into the center of a cylinder whose walls have been coated with mineral oil. As they fall, the animals fly out from the center of the cylinder, strike the cylinder walls, and are trapped in the oil. In practice, the animals remain at the position where they first collide with the wall. Animals that fly well tend to accumulate near the top of the column, whereas animals that fly poorly are predominately found near the bottom of the column.

To determine whether the defective flight phenotype of *slowpoke* mutants was associated with a muscle defect, we assayed the flight ability of $ash2^{18}/slo^4$ transheterozygotes. In this assay, these animals accumulated near the top of the column and at the same position as wild-type flies. As shown in Figure 5, the slo^4 homozygotes were obviously impaired in their flying ability and accumulated in the bottom half of the column.

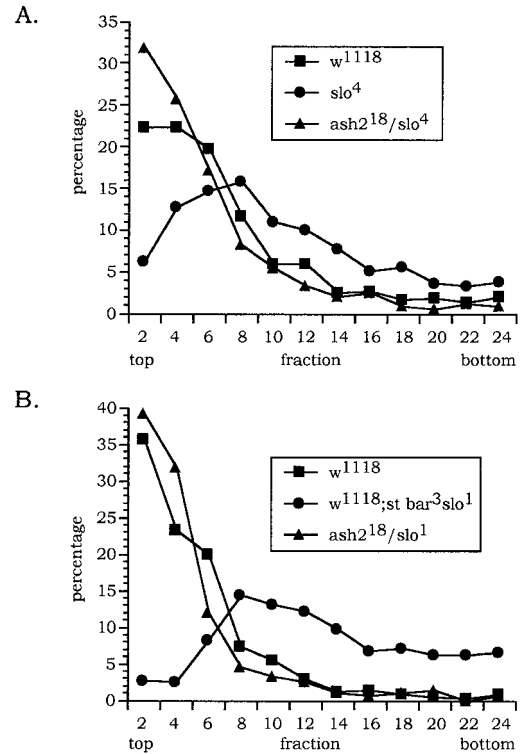


Figure 5. Column-based flight assay used to measure the relative capacity of flies for flight. Flies are dropped from a vial into the center of an oil-coated cylinder. The falling flies fly from the center and are trapped in the oil. The distance that they fall is correlated with their capacity for flight (Benzer, 1973; Elkins et al., 1986; Green et al., 1986). The column was fractionated, top to bottom, into bins 5 cm in length. The abscissa represents the distance from the top of the column that the flies fell. The ordinate is the percentage of animals assayed. *A*, The $ash2^{18}$ chromosome was tested for its capacity to complement the flight defect associated with the slo^4 mutant allele. Results from an assay performed using 952 w^{1118} , 828 slo^4 , and 733 $ash2^{18}/slo^4$ transheterozygotes. *B*, Ability of the $ash2^{18}$ chromosome to complement the flight defect associated with the slo^1 mutant allele. Results from an assay performed using 1002 w^{1118} , 903 $w^{1118}; st bar^3 slo^1$, and 369 $ash2^{18}/st bar^3 slo^1$ flies. The w^{1118} stock carries a wild-type copy of the *slowpoke* gene and served as the positive control. Wild-type flies (*squares*) and the $ash2^{18}/slo^4$ or $ash2^{18}/slo^1$ transheterozygotes (*triangles*) accumulate near the top of the column. The slo^4 or slo^1 homozygotes (*circles*) accumulate deeper in the column. Flies that did not initiate flight are not counted and are trapped in a pool of oil at the bottom of the column.

Sticky-feet phenotype

After a brief heat pulse from 22 to 37°C, flies that are homozygous for null mutations in the *slowpoke* gene have been described as standing motionless for several minutes (Elkins et al., 1986). This behavior is better described as a sticky-feet phenotype. Approximately 15 sec after a 2–8 min, 37–40°C heat pulse, the flies stand in place and can be pushed with a pencil. During this time, they behave as if their feet are stuck to the surface on which they stand (Fig. 6). Continuing to push the fly causes it to gradually lean over and eventually to fall onto its side or back. The flies are not paralyzed per se, because once knocked over they right themselves, after which, their feet frequently stick to the surface. If the fly is left undisturbed, this behavior can persist for many minutes. Recovery seems to be speeded by repeated touching of the fly. All of the known *slowpoke* null alleles ($slo^1, slo^2, slo^3, slo^4, slo^5$, and slo^8) exhibit this behavior as homozygotes and in all allelic

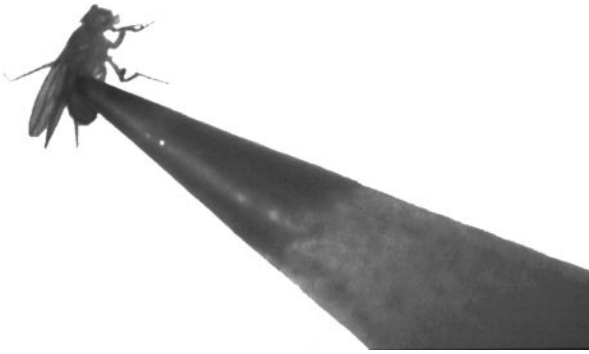


Figure 6. An example of the sticky-feet phenotype exhibited by animals homozygous for null mutations in the *slowpoke* gene. This particular homozygous *slo*⁴ male has been exposed to a bright light (see Materials and Methods) for 15 sec and is being pushed with a number 2 pencil. Such animals do not attempt to escape or avoid the pencil and hang onto the surface on which they stand. The animal is shown leaning over in response to the pressure. With continued pressure, the animal will topple over and then in a very uncoordinated manner attempt to right himself. If successful, such an animal will usually once again exhibit the sticky-feet phenotype. Recovery from the behavior can take many minutes and seems to be speeded by repetitive stimulation.

combinations. Heterozygous and wild-type animals do not manifest this response.

We have also been able to elicit this behavior using a cool but very bright light. Stimulation with light evokes the behavior much more rapidly than exposure to an elevated temperature. Homozygous *slo*¹ flies exposed to the maximal output from a 60 W fiber optic lamp elicit this behavior in 15 sec. During this time, no change in temperature of the platform was observed. Based on this result, we postulate that the sticky-feet behavior is induced not by temperature but by overstimulation of the animals.

The *ash2*¹⁸ deletion removes promoters C0 and C1 (neural-specific) but leaves promoter C2 intact (muscle and tracheal cell-specific). This provides us with the opportunity to determine whether the sticky-feet behavior arises from a lack of expression in muscle or neural tissue. Flies carrying a single copy of the *ash2*¹⁸ and *slo*⁴ mutant chromosomes were behaviorally tested. Animals that walked or flew away when pushed on with a pencil were scored as exhibiting wild-type behavior.

Flies carrying a single copy of the *ash2*¹⁸ and *slo*⁴ mutant chromosomes were produced by crossing *yw*; *ash2*¹⁸, *red*, *ely*⁺ TM3 *Sb Ser* virgin females to *slo*⁴ males. The genotypes of these animals were (1) *yw*; *ash2*¹⁸/*slo*⁴ male, (2) *yw*; *slo*⁴/*ly*⁺, TM3 *Sb Ser* male, (3) *yw*/+; *ash2*¹⁸/*slo*⁴ female, and (4) *yw*/+; *slo*⁴/*ly*⁺ TM3 *Sb Ser* female. Although all of the progeny were behaviorally scored, the data reported here was collected only from females. The males were excluded because behavioral testing of the males could not be performed as a blind assay because the *ash2*¹⁸/*slo*⁴ transheterozygote and *slo*⁴ heterozygote males can be distinguished with the naked eye based on their body color. However, detection of the physical markers that distinguish the transheterozygote and heterozygote females require substantial magnification and therefore permit an unbiased behavioral assay to be performed.

A total of 516 females were examined. None of the 238 *yw*/+; *slo*⁴/*ly*⁺ TM3 *Sb Ser* animals showed the sticky-feet phenotype and were behaviorally scored as wild type. Of the remaining *yw*/+; *ash2*¹⁸/*slo*⁴ animals, 254 were scored as exhibiting the sticky-feet phenotype, and 24 were scored as exhibiting wild-type

behavior. As a control, we performed the behavioral assay on 258 *slo*⁴ animals. Of these, 230 were scored as having the sticky-feet phenotype and 28 were scored as wild type. It is common for us to have a 10% mis-scoring rate for the *slo*⁴ homozygous parental stock. Therefore, we conclude that the *ash2*¹⁸ mutation fails to complement this *slowpoke* behavioral phenotype.

DISCUSSION

Mutations in the *slowpoke* gene have pleiotropic effects on animal physiology and behavior. In both muscles and neurons, *slowpoke* mutations have been shown to eliminate the BK-type Ca-activated K current called *I*_{CF} (Salkoff, 1983; Elkins et al., 1986; Gho and Mallart, 1986; Singh and Wu, 1989; Komatsu et al., 1990; Saito and Wu, 1991; Broadie and Bate, 1993). In muscles, a *slowpoke* null allele alters the shape and duration of the flight muscle action potentials (Elkins et al., 1986; Elkins and Ganetzky, 1988). In neurons, the same mutation affects not only the action potentials shape (Saito and Wu, 1991) but also the release of neurotransmitter from motoneurons (Gho and Ganetzky, 1992; Warbington et al., 1996) and the habituation of a neuronal circuit in the adult CNS (Engel and Wu, 1998). This indicates that *slowpoke* channels are of central importance for the normal function of both neurons and muscle fibers.

During repetitive stimulation of the flight muscle, wild-type animals produce trains of well formed action potentials. Animals homozygous for a *slowpoke* null mutation produce extremely broad action potential spikes. In a train of action potentials, the first spike is typically of normal breadth. Subsequent spikes, however, are typically extremely broad and often have two peaks. In *slowpoke* mutants, it is believed that a current called *I*_A, conducted by the *Shaker*-encoded voltage-gated K channel, is responsible for repolarization of the first spike (Elkins and Ganetzky, 1988). The voltage-dependent inactivation of this current reduces its contribution to the repolarization of subsequent spikes. In wild-type muscle, the *slowpoke*-encoded channels ensure the rapid repolarization of subsequent spikes. However, in muscle lacking a functional *slowpoke* gene, the subsequent spikes cannot be properly repolarized.

Animals carrying null mutations in the *slowpoke* gene also present behavioral phenotypes. Although the animals are very healthy and fecund, they are relatively lethargic and have a limited capacity for flight. They also manifest a temperature- or light-induced sticky-feet phenotype. *A priori*, one cannot predict whether these behavioral traits have a neuronal or muscular origin. Previous studies of the *slowpoke* transcriptional control region indicates that neuronal and muscle expression arise from different promoters, which are separated by >3.7 kb of genomic DNA (Brenner and Atkinson, 1996; Brenner et al., 1996). An ideal tool for identifying the origin of the sticky-feet and flight phenotypes would be a mutant lesion that affected either the neuronal or muscle promoters but not both. The *ash2*¹⁸ deletion provides just such a tool. This deletion removes the neighboring *ash2* gene and the neuronal promoters of the *slowpoke* gene. The portion that remains intact includes promoter C2 and other sequences required for muscle expression (Brenner and Atkinson, 1996; Brenner et al., 1996). Therefore, the *ash2*¹⁸ deletion should also be viewed as a *slowpoke* mutation that eliminates neuronal *slowpoke* expression. In addition, because the *ash2*¹⁸ deletion does not involve the *slowpoke* coding region, the BK channels expressed by the chromosome should be fully functional.

As an aside, the *ash2*¹⁸ deletion also allows us to determine the

orientation of the *slowpoke* gene on *Drosophila* chromosome 3. Genetic mapping indicates that *slowpoke* (genetic position 3–86) is distal to *ash2* (genetic map position 3–78.6). That is, the *slowpoke* locus is farther from the centromere than the *ash2* locus. The *ash2*¹⁸ deletion removes both *ash2* and a portion of the *slowpoke* transcription control region but not the *slowpoke* coding region. Therefore, the 5' end of *slowpoke* transcription unit must be closer to *ash2* than the 3' end of the transcription unit, which means that *slowpoke* is positioned on chromosome 3 such that transcription proceeds away from the centromere.

As predicted, the *ash2*¹⁸ chromosome complements the *slo*⁴ mutant muscle phenotype with regard to its electrophysiological abnormality. Furthermore, the restoration of electrical properties is correlated with a restoration of normal flight. This strongly suggests that the flight defect in animals carrying *slowpoke* null mutations is caused solely by an absence of BK-type channels in muscle and the resultant abnormalities in the electrical character of the muscle fiber. Satisfyingly, a phenotype suspected to be neuronal in origin is not complemented by the *ash2*¹⁸ chromosome; that is, the *ash2*¹⁸/*slo*⁴ transheterozygotes exhibit a robust sticky-feet phenotype.

The sticky-feet phenotype is an extremely unusual behavior. It is triggered by both heat and bright light, suggesting that it is a direct response to overstimulation of the animal. The persistence of the behavior for many minutes after the stimulus ends suggests that the affected cells enter a prolonged state of inappropriate activity. Such a response might be caused by seizure-like activity within a circuit involved in evoking a reflex behavior.

REFERENCES

- Adamson AL, Shearn A (1996) Molecular genetic analysis of *Drosophila ash2*, a member of the trithorax group required for imaginal disc pattern formation. *Genetics* 144:621–633.
- Adelman JP, Shen KZ, Kavanaugh MP, Warren RA, Wu YN, Lagrutta A, Bond CT, North RA (1992) Calcium-activated potassium channels expressed from cloned complementary DNAs. *Neuron* 9:209–216.
- Atkinson NS, Robertson GA, Ganetzky B (1991) A component of calcium-activated potassium channels encoded by the *Drosophila slo* locus. *Science* 253:551–555.
- Baumann A, Krah-Jentgens I, Müller R, Müller-Holtkamp F, Seidel R, Kecskemethy N, Casal J, Ferrus A, Pongs O (1987) Molecular organization of the maternal effect region of the *Shaker* complex of *Drosophila*: characterization of an I_A channel transcript with homology to vertebrate Na⁺ channel. *EMBO J* 6:3419–3429.
- Becker MN, Brenner R, Atkinson NS (1995) Tissue-specific expression of a *Drosophila* calcium-activated potassium channel. *J Neurosci* 15:6250–6259.
- Benzer S (1973) Genetic dissection of behavior. *Sci Am* 229:24–37.
- Bohm RA, Brenner R, Wang B, Atkinson N (2000) Transcriptional control of Ca-activated K channel expression: identification of a second, evolutionarily-conserved, neuronal promoter. *J Exp Biol* 203:693–704.
- Brenner R, Atkinson N (1996) Developmental and eye-specific transcriptional control elements in an intronic region of a Ca²⁺-activated K⁺ channel gene. *Dev Biol* 177:536–543.
- Brenner R, Atkinson NS (1997) Calcium-activated potassium channel gene expression in the midgut of *Drosophila*. *Comp Biochem Physiol B Biochem Mol Biol* 118:411–420.
- Brenner R, Thomas TO, Becker MN, Atkinson NS (1996) Tissue-specific expression of a Ca²⁺-activated K⁺ channel is controlled by multiple upstream regulatory elements. *J Neurosci* 16:1827–1835.
- Broadie KS, Bate M (1993) Development of larval muscle properties in the embryonic myotubes of *Drosophila melanogaster*. *J Neurosci* 13:167–180.
- Elkins T, Ganetzky B (1988) The roles of potassium currents in *Drosophila* flight muscles. *J Neurosci* 8:428–434.
- Elkins T, Ganetzky B, Wu C-F (1986) A *Drosophila* mutation that eliminates a calcium-dependent potassium current. *Proc Natl Acad Sci USA* 83:8415–8419.
- Engel JE, Wu C-F (1992) Interactions of membrane excitability mutations affecting potassium and sodium currents in the flight and giant fiber escape systems of *Drosophila*. *J Comp Physiol [A]* 171:93–104.
- Engel JE, Wu C-F (1998) Genetic dissection of functional contributions of specific potassium channel subunits in habituation of an escape circuit in *Drosophila*. *J Neurosci* 18:2254–2267.
- Gho M, Ganetzky B (1992) Analysis of repolarization of presynaptic motor terminals in *Drosophila* larvae using potassium-channel-blocking drugs and mutations. *J Exp Biol* 170:93–111.
- Gho M, Mallart A (1986) Two distinct calcium-activated potassium currents in larval muscle fibres of *Drosophila melanogaster*. *Pflügers Arch* 407:526–533.
- Green CC, Sparrow JC, Ball E (1986) Flight testing columns. *Drosophila Information Service* 63:141.
- Kamb A, Iverson LE, Tanouye MA (1987) Molecular characterization of *Shaker*, a *Drosophila* gene that encodes a potassium channel. *Cell* 50:405–413.
- Komatsu A, Singh S, Rathe P, Wu C-F (1990) Mutational and gene dosage analysis of calcium-activated potassium channels in *Drosophila*: correlation of micro- and macroscopic currents. *Neuron* 4:313–321.
- Pacha J, Frindt G, Sackin H, Palmer LG (1991) Apical maxi K channels in intercalated cells of CCT. *Am J Physiol* 261:F696–F705.
- Papazian DM, Schwarz TL, Tempel BL, Jan YN, Jan LY (1987) Cloning of genomic and complementary DNA from *Shaker* a putative potassium channel gene from *Drosophila*. *Science* 238:749–753.
- Saito M, Wu C-F (1991) Expression of ion channels and mutational effects in giant *Drosophila* neurons differentiated from cell division-arrested embryonic neuroblasts. *J Neurosci* 11:2135–2150.
- Salkoff L (1983) *Drosophila* mutants reveal two components of fast outward current. *Nature* 302:249–251.
- Salkoff LB, Wyman RJ (1983) Ion currents in *Drosophila* flight muscles. *J Physiol (Lond)* 337:687–709.
- Singh A, Singh S (1999) Unmasking of a novel potassium current in *Drosophila* by a mutation and drugs. *J Neurosci* 19:6838–6843.
- Singh S, Wu C-F (1989) Complete separation of four potassium currents in *Drosophila*. *Neuron* 2:1325–1329.
- Stoner LC, Morley GE (1995) Effect of basolateral or apical hyposmolarity on apical maxi K channels of everted rat collecting tubule. *Am J Physiol* 268:F569–F580.
- Tsunoda S, Salkoff L (1995) The major delayed rectifier in both *Drosophila* neurons and muscle is encoded by *Shab*. *J Neurosci* 15:5209–5221.
- Warbington L, Hillman T, Adams C, Stern M (1996) Reduced transmitter release conferred by mutations in the *slowpoke*-encoded Ca²⁺-activated K⁺ channel gene of *Drosophila*. *Invert Neurosci* 2:51–60.
- Wei A, Jegla T, Salkoff L (1996) Eight potassium channel families revealed by the *C. elegans* genome project. *Neuropharmacology* 35:805–829.