

## Energetics of Gliding Motility in *Mycoplasma mobile*

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***Mycoplasma mobile* glides on surfaces at up to 7  $\mu\text{m/s}$  by an unknown mechanism. We studied the energetics that power gliding by using a novel, growth medium-free system. We found that cells could glide in defined media if the glass substrate is preconditioned by exposure to horse serum. The active component that potentiates gliding is sensitive to proteinase K treatment. We used the defined medium system to test the effect of various inhibitors, ionophores, and poisons on motility of *M. mobile*. Valinomycin, carbonyl cyanide 4-(trifluoromethoxy)phenylhydrazone (FCCP), *N,N'*-dicyclohexylcarbodiimide, phenamil, amiloride, rifampin, and puromycin had no short-term effects on gliding. We also confirmed that we were able to modulate the membrane potential with valinomycin and FCCP by using a potential-sensitive dye. Shifting the pH likewise had no effect on motility. These results rule out the use of conventional ion motive forces to power gliding. Arsenate had a dramatic inhibitory effect on gliding, and both the speed and the fraction of cells moving tracked ATP levels. Sodium orthovanadate had a slight but significant inhibitory effect on gliding. Taken together, these results suggest that the motor system of *M. mobile* is likely an ATPase or is directly coupled to an ATPase.**

While studies of locomotion in swimming bacteria are well advanced, investigations of gliding motility remain comparatively limited. Yet gliding motility, defined as a smooth translocation over a solid surface, is represented frequently throughout the eubacterial phylogenetic tree and in some instances has been associated with pathogenicity (25). Even several species of mycoplasmas, some of the simplest bacteria known in terms of size and genomic content, are known to perform gliding motility (14, 32).

The mycoplasmas are wall-less bacteria characterized by small physical dimensions and genome sizes (32). Among the mycoplasmas, the fish pathogen *Mycoplasma mobile* demonstrates extremely robust gliding motility (16, 34). *M. mobile* is one of the flask-shaped mycoplasmas (approximately  $1.0 \times 0.3 \mu\text{m}$ ) and has a genome of approximately 780 kbp (4). It has always been observed to glide in the direction of the “head” (corresponding to the tapered end of the cell) without reversals or pauses at speeds of up to 7  $\mu\text{m/s}$  (34). It can tow an erythrocyte, roughly 10 times its size, without significant loss in speed and has been measured to exert up to 27 pN of force (28, 33). Some recent progress at uncovering the molecular mechanism of gliding in *M. mobile* has been made, including localization of the gliding apparatus to the head region of its flask-like cell body and isolation of mutants with altered gliding phenotypes (29, 30, 41). However, little is known about the prerequisites or energy source for gliding in *M. mobile*.

Flagellated bacteria use ion motive forces to power their flagellar motors. For instance, *Escherichia coli* uses a proton motive force ( $\Delta p_{\text{H}}$ ), while various *Vibrio* species use a sodium

motive force ( $\Delta p_{\text{Na}}$ ) to drive their polar (but not lateral) flagella (3, 18, 19, 24). The energy source for motility has been investigated in some genera of gliding bacteria as well. In the case of the *Flavobacteria*, gliding motility appears to be powered by  $\Delta p_{\text{H}}$ , while the social motility system of the *Myxobacteria* relies on type IV pili and, therefore, ATP hydrolysis (25).

The mycoplasmas seem to lack any form of respiration and generate ATP through fermentation of sugars and substrate-level phosphorylation (32). It is known that mycoplasmas can generate a transmembrane potential ( $\Delta\Psi$ ) ranging from  $-28$  to  $-48$  mV (negative inside the cell) and a  $\Delta p_{\text{H}}$  ranging from  $-52$  to  $-72$  mV (37). Therefore, it would seem that mycoplasmas could use either an ion motive force or direct utilization of ATP to power motility. We set out to develop a medium-free system in which we could study gliding of *M. mobile* on glass and determine its energy source.

### MATERIALS AND METHODS

**Reagents.** Heart infusion broth and yeast extract were from Becton Dickinson (Sparks, Md.). 3,3'-dipropylthiadicarbocyanine iodide ( $\text{DiSC}_3[5]$ ) was from Molecular Probes (Eugene, Oreg.). The ENLITEN ATP measurement system was from Promega (Madison, Wis). All other reagents were from Sigma-Aldrich (St. Louis, Mo.). Water was 18 M $\Omega$  deionized ( $\text{dH}_2\text{O}$ ).

**Strains.** *M. mobile* strain 163K (ATCC 43663) was grown to an optical density at 600 nm ( $\text{OD}_{600}$ ) of 0.07 to 0.10 in plastic tissue culture flasks at 22°C in Aluotto medium consisting of 10% inactivated horse serum, 2.1% beef heart infusion broth, and 0.56% yeast extract adjusted to pH 7.8 and supplemented with 50 mg of ampicillin/liter and 250 mg of thallium acetate/liter (1).

**Preparation of coverslips.** Circular glass coverslips were subjected to the following sequence of treatments (all at room temperature with gentle agitation): 10 min in saturated ethanolic KOH, four 5-min changes in  $\text{dH}_2\text{O}$ , 15 min in inactivated horse serum, and three 5-min changes in  $\text{dH}_2\text{O}$ . The coverslips were then left to dry in a laminar flow hood and stored at room temperature until use, resulting in a preparation that was stable for at least 4 weeks. Note that fetal bovine serum can also be used with equal effectiveness.

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**Protease treatment.** Prepared coverslips were digested overnight with 20 mg of proteinase K/ml (or dH<sub>2</sub>O as a control) at 42°C in a humid environment and washed with four 5-min changes in dH<sub>2</sub>O.

**Buffers.** The following buffers were used: phosphate-buffered saline (PBS; 150 mM NaCl, 50 mM sodium phosphate [pH 8.0]), PBS/G (PBS [pH 8.0] plus 10 mM glucose), PBS-K/G (140 mM NaCl, 10 mM KCl, 50 mM sodium phosphate, pH 8.0 [or other pH as specified], 10 mM glucose), ArBS-K/G (140 mM NaCl, 7.5 mM KCl, 47.5 mM sodium arsenate, 2.5 mM potassium arsenate [pH 8.0], 10 mM glucose), and valinomycin buffer (100 mM NaCl, 50 mM KCl, 50 mM sodium phosphate [pH 8.0], 10 mM glucose).

**Motility assay.** Comparisons were made of gliding speeds of cells in a given buffer and cells in the same buffer containing the compound to be tested, referred to as control buffer and test buffer, respectively. Cells (diluted to an OD<sub>600</sub> of 0.025 in fresh medium [1 ml]) were centrifuged at room temperature for 10 min at 10,000 × g, washed in 1 ml of control buffer or test buffer, and centrifuged again in the same manner. The final pellet was suspended in 50 μl of the control buffer or test buffer. Six microliters of this suspension was pipetted onto a prepared coverslip and incubated at room temperature for 15 min in a humid environment. During that time, a flow chamber (5) was assembled with a spare coverslip ringed with Apiezon-L grease (GEC-Alsthom Ltd., Manchester, United Kingdom) and equilibrated by continuous flow with control or test buffer for at least 10 min. Coverslips with adhered cells were then ringed with grease and inverted onto the flow chamber. All observations were made at room temperature (~22°C). Control buffer or test buffer was drawn into the flow chamber at the rate of 100 μl/min for 10 min to remove nonadherent cells, and then the motion of the remaining cells was recorded (with the flow left on). Cells were visualized with a Nikon Optiphot microscope equipped with a 60× oil immersion phase-contrast objective, a Zeiss optovar set to 1.25×, and a 10× projection lens. Recordings were made with a charge-coupled device camera connected to a Mini-DV recorder (Sony GV-D1000, New York, N.Y.). In the standard assay, 10-s recordings were made every 10 min over the course of 30 min. Time zero was defined to occur at the time of the first recording. Video was captured to a computer with Adobe (San Jose, Calif.) Premier software at a rate of 10 frames/s. The speed of gliding mycoplasmas was computed for frames 40 to 59 of each 10-s (100-frame) time point by using a particle tracking program written by Darnton et al. (8). Slight modifications were made to Darnton's program to output the proper statistics (the MATLAB scripts are available at <http://www.rowland.org/labs/bacteria/index.html>). Speeds of 30 to 50 cells were measured at each time point in each experiment. The compound being tested was considered to have a significant effect on motility only if a comparison of the test and control populations by a two-tailed Student's *t* test had a *P* value of <0.05 at all time points studied.

**pH shift.** Cells were prepared in PBS-K/G (pH 8.0) and then shifted to PBS-K/G at the desired pH after the first recording was taken. The second recording was taken at *t* = 5 min.

**Arsenate and ATP.** Measurements of motility parameters and ATP levels were made in parallel by slightly modifying the standard motility assay. Eleven 1-ml aliquots of cells at an OD<sub>600</sub> of 0.025 were washed in PBS-K/G as described above. Cells in one tube were resuspended in 50 μl of PBS-K/G and used for the flow chamber experiments where they were treated as described above, while the remaining 10 tubes were washed an additional time in PBS-K/G to simulate the wash experienced by the cells in the flow chamber. At time zero of the experiment, the buffer feeding the flow chamber was either switched to ArBS-K/G or allowed to remain in PBS-K/G in the case of the control. Concurrently, each of the remaining 10 tubes was resuspended in 100 μl of ArBS-K/G or PBS-K/G, depending on the experiment. Ten-second recordings were made every 10 min as was done before, but the time course was extended to 90 min because there was no preincubation with test buffer. At each time point, one tube of cells was extracted for ATP by adding 2 μl of trichloroacetic acid to the tube and vortexing and then adding 12 μl of 1.5 M Tris base to neutralize the acid (12). The sample was then diluted with 342 μl of 5 mM Tris buffer (pH 7.75) and frozen at -20°C until the ATP assay was performed. In some cases, further dilution of the sample with 5 mM Tris (pH 7.75) was necessary to get the sample within the range of the assay standard curve. For the ATP assay, 10 μl of each sample was added to a glass vial, and then 100 μl of ENLITEN rL/L reagent was added to the tube and mixed by repeated pipetting. Measurements were taken exactly 15 s after the addition of the rL/L reagent in a home-built luminometer consisting of a photomultiplier tube connected to a current-to-voltage converter and a chart recorder (27). Two measurements were made for each sample and were compared to a standard curve of ATP constructed from duplicated measurements of samples of ATP ranging from 31.3 to 500 fmol.

**Membrane potential.** Cells were grown as described above, and a sufficient amount of culture was centrifuged at 10,000 × g such that resuspension in 2 ml

of test buffer resulted in an OD<sub>600</sub> of 0.175 to 0.200. This suspension was placed in a cuvette, and DiS-C<sub>3</sub>-[5] was added from an EtOH stock to a final concentration of 150 nM (final EtOH, 0.05%). Fluorescence was monitored at excitation and emission wavelengths of 625 and 667 nm, respectively, in a Fluoromax fluorimeter (Spex, Edison, N.J.) equipped with a magnetic stirring cell. The dye was allowed to partition into the cells until a stable signal was obtained (4 to 5 min), and then the test substance (valinomycin, carbonyl cyanide 4-(trifluoromethoxy)phenylhydrazone [FCCP], or *N,N'*-dicyclohexylcarbodiimide [DCCD]) was added. Changes in the fluorescence signal were observed within a few seconds of the addition of the test substance if changes were observed at all.

**Isolation of proteins that bind to glass from horse serum.** Approximately 100 μl of glass beads (425 to 600 μm in diameter) were prepared in exactly the same way as the coverslips described above. After drying, the beads were boiled in Laemmli sodium dodecyl sulfate-polyacrylamide gel electrophoresis buffer (36). The resulting sample was loaded to a sodium dodecyl sulfate-4 to 12% polyacrylamide gel electrophoresis gradient gel (Invitrogen, Carlsbad, Calif.) and subjected to electrophoresis. Major bands were excised and subjected to in-gel tryptic digestion and identification by ion trap mass spectrometry (13).

## RESULTS

**Conditioning of glass.** In pilot experiments for this project, we noticed that *M. mobile* cells would glide on glass surfaces if they were pelleted by centrifugation and then simply resuspended in a pure buffer test medium (e.g., PBS/G). However, when an intervening wash step was included, cells would adhere to glass in aggregates but would not glide. Evidently, in the former case, an essential component from the medium that potentiates gliding was carried over from the pellet, while in the latter case, the extra wash step eliminated the component. We tested the major components of the growth medium (horse serum, beef heart infusion, and yeast extract) by adding each one back to a washed culture and found that only horse serum restored motility. We subsequently found that a simple pretreatment of glass with fresh horse serum was enough to potentiate the motility of washed cells. Therefore, the factor that potentiates gliding on glass surfaces is native to horse serum and is not a factor produced by mycoplasmas in culture. Subsequent to this discovery, we adopted the coverslip pretreatment method described in Materials and Methods.

**Identity of conditioning factor.** We subjected pretreated coverslips to overnight incubation with either proteinase K or water. The coverslips were then washed with water to eliminate the protease so that gliding could be tested. We found that coverslips treated overnight with proteinase K no longer facilitated gliding motility of a washed culture, whereas the control coverslips treated with water alone still potentiated gliding of mycoplasmas. However, the coverslips treated with proteinase K still allowed adherence of cells to the glass with approximately the same number of cells stuck as is found in the control. We therefore conclude that a protein or combination of proteins either directly or indirectly mediates the ability of *M. mobile* cells to glide on glass surfaces.

As albumin is the most abundant protein known in serum, we pretreated coverslips as described above using a solution of bovine serum albumin (1 mg/ml) in place of the horse serum; note that the pretreatment of coverslips with fetal bovine serum also potentiated gliding. This type of pretreatment did not potentiate gliding of washed cells, so a protein or proteins other than albumin are probably involved in conditioning the glass surface. We attempted to isolate this component(s) by identifying the major species that bound to glass from horse serum. Mass spectrometry was able to identify albumin (as

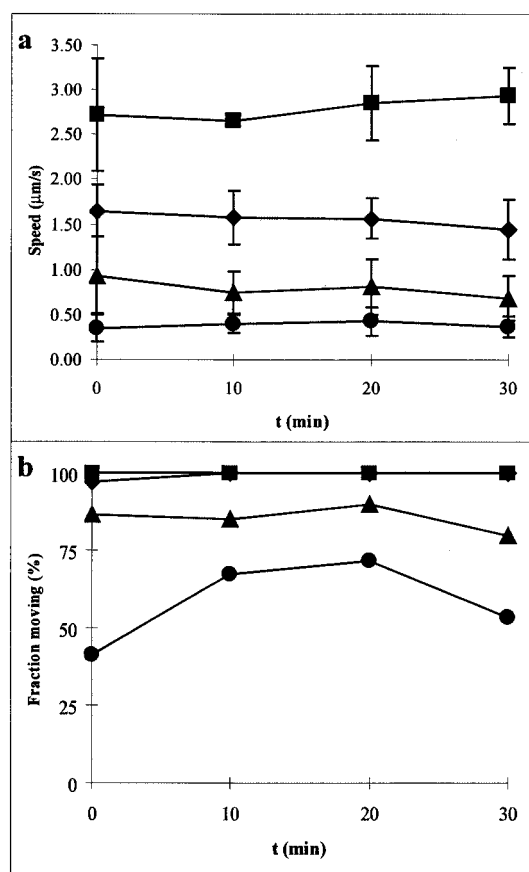


FIG. 1. Gliding in growth and artificial media. (a) Gliding speed of motile cells. Only cells moving at speeds greater than  $0.25 \mu\text{m/s}$  are considered. Error bars represent one standard deviation. (b) Percentage of cells moving at speeds greater than  $0.25 \mu\text{m/s}$ . Symbols: ■, growth medium; ◆, PBS-K/G; ▲, PBS/G; ●, PBS.

expected), immunoglobulin, and casein as the top three proteins bound to glass beads. However, attempts to coat glass coverslips with solutions of these proteins (1 mg/ml each) did not potentiate motility. Rather, immunoglobulin and casein abolished cytoadherence of the cells, while albumin caused clumps of nonmotile cells to adhere to the glass surface. The actions of these proteins in combination have not yet been tested.

**Gliding of *M. mobile* in artificial media.** The development of the pretreatment procedure for glass coverslips allowed us to study motility in artificial media (e.g., simple buffer systems). Movies of *M. mobile* gliding can be seen at [http://www.rowland.org/labs/bacteria/showmovie.php?mov=mycoplasma\\_mobile](http://www.rowland.org/labs/bacteria/showmovie.php?mov=mycoplasma_mobile). A comparison of the basic buffers used in subsequent experiments is shown in Fig. 1. While none of the artificial media used resulted in gliding speeds as rapid as those in growth medium (Fig. 1a), most media supported motility of  $>80\%$  of the cells (Fig. 1b). The omission of glucose from the medium caused a substantial decrease in both speed and percentage of cells that were moving. The sole source of ATP generation in most mycoplasmas is through substrate-level phosphorylation during glycolysis, although some mycoplasmas can use arginine as an energy source (32). This indicates that ATP synthesis may play a key role in the

energetics of mycoplasma motility. We found that sucrose could also be used in place of glucose as an energy source in motility experiments (data not shown), indicating the ability of *M. mobile* to metabolize sucrose.

Gliding speeds of mycoplasma in medium containing  $\text{Na}^+$  and glucose (PBS/G) were consistently around  $1 \mu\text{m/s}$ , but the cells appeared thin and elongated. The inclusion of a small amount of  $\text{K}^+$  (10 mM KCl) in the medium caused a 33 to 50% increase in gliding speed, and the cells looked more like those in growth medium. However, the addition of too much  $\text{K}^+$  ( $>100 \text{ mM}$ ) caused cells to swell and become coccus shaped, which inhibited motility but not glass binding (data not shown). Mycoplasmas are known to selectively accumulate  $\text{K}^+$  to over 200 mM against a gradient (37). This fact, combined with these observations on gliding in artificial media of different ionic compositions, suggests that  $\text{K}^+$  plays an important role in osmoregulation and shape determination in *M. mobile* and, in turn, that osmoregulation and shape determination are important factors in gliding motility.

We also tested the ability of cells to glide in a medium in which almost all cations were completely impermeant to the membrane (200 mM choline chloride,  $200 \mu\text{M}$  HEPES [pH 7.9], 10 mM glucose, and final  $\text{Na}^+$  of about  $160 \mu\text{M}$  due to titration with NaOH). We found that  $>80\%$  of cells glided in this medium, achieving speeds of up to  $2.6 \pm 0.46 \mu\text{m/s}$ . However, gliding cells tended to fall off of the glass surface after a short period of observation (10 min) for reasons that are not yet understood.

**Effects of external pH.** If a proton motive force is involved in gliding, changing the  $\Delta\text{pH}$  might have an effect on motility. We prepared all cells in pH 8.0 buffer and then subsequently shifted them to a lower pH in the flow chamber. Shifting the external pH of the medium did not have a significant effect on speed or percentage of motile cells (Fig. 2).

**Effects of ionophores, inhibitors, and poisons.** We examined the effects of several compounds known to interfere with various cellular processes that have been shown to be involved in the motility of other organisms. In all cases, motility and the percentage of motile cells were observed over a 30-min period and compared to a carefully matched buffer control that did not contain the compound in question. Buffer formulations are listed in Table 1.

The addition of the potassium ionophore valinomycin to cells in the presence of high external  $\text{K}^+$  would be expected to clamp  $\Delta\Psi$  at a lower-than-normal value (37, 38). Valinomycin had no significant effect on motility compared to the control (Fig. 3a and b), even in the presence of 50 mM external potassium. This result was interesting, considering that high amounts of potassium caused cells to become coccoid and stop gliding (see above). However, at the end of the 30-min observation period, nearly all cells were still moving at a rate comparable to that of the matched control. Similarly, the protonophore FCCP did not cause a significant decrease in either gliding speed or percentage of cells that were gliding (Fig. 3c and d). FCCP is reported to abolish  $\Delta\text{pH}$  in various mycoplasmas (37), and the inclusion of FCCP in the medium would be expected to short circuit any proton-driven motor by providing an alternate conduction path for protons and driving  $\Delta\text{pH}$  toward 0.

We studied the effect of valinomycin and FCCP on the



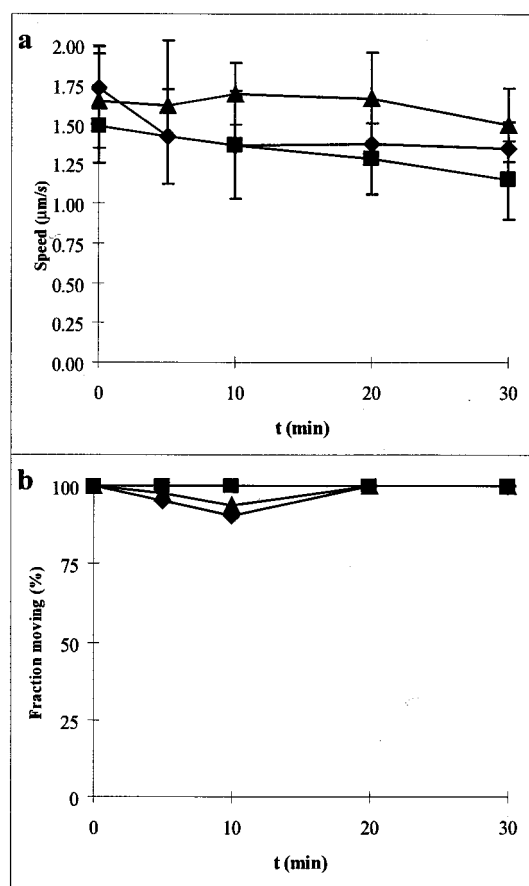


FIG. 2. Effect of shifting pH on gliding motility. All samples were prepared in PBS-K/G (pH 8.0) and then shifted to the indicated pH at  $t = 1$  min. (a) Gliding velocity; (b) fraction of cells moving. Symbols: ■, pH 8; ◆, pH 7; ▲, pH 6.

membrane potential ( $\Delta\Psi$ ) of *M. mobile* by using the positively charged fluorescent indicator DiSC<sub>3</sub>[5]. This compound partitions between the cell and the external medium at higher intracellular concentrations when the inside of the cell is more negative. At higher concentrations, the dye dimerizes or aggregates, quenching the fluorescence (44, 45). Thus, a decrease in fluorescence indicates membrane hyperpolarization. Under the conditions of the gliding experiment, the addition of valinomycin to the cells caused a depolarization of the membrane,

while the addition of FCCP caused a slight hyperpolarization of the membrane (Fig. 4a). These shifts occurred within seconds after addition of either compound, so their effects must have been manifested during the gliding experiments.

Mycoplasmas are expected to have an internal  $K^+$  of about 200 mM (37). The addition of valinomycin in the presence of 50 mM external  $K^+$  (the concentration used in the gliding experiment) would be expected to clamp  $\Delta\Psi$  at  $RT/F \ln(50 \text{ mM } K^+_{\text{ext}}/200 \text{ mM } K^+_{\text{int}}) = -35 \text{ mV}$ , where  $R$  is the gas constant,  $T$  is the absolute temperature, and  $F$  is the Faraday. We believe that the normal  $\Delta\Psi$  of *M. mobile* is greater than  $-76 \text{ mV}$  because an analogous experiment that we performed with 10 mM external  $K^+$  still induced depolarization upon the addition of valinomycin. Thus, in the gliding experiment,  $\Delta\Psi$  was reduced by more than a factor of two.

The experiment also shows that FCCP affects  $\Delta\Psi$  as well as  $\Delta\text{pH}$  (37). The sign and magnitude of the shift in  $\Delta\Psi$  depends on the external pH. At a higher pH (pH 7.0 or 8.0), FCCP caused a hyperpolarization, while at a lower pH (pH 6.0), FCCP caused a depolarization (Fig. 4b). Gliding motility was not affected by external pH in this range (Fig. 2). Therefore, by employing valinomycin and FCCP, we were able to collapse both  $\Delta\Psi$  and  $\Delta\text{pH}$  (and therefore modulate  $\Delta\text{pH}$ ) without significant effect on gliding motility. Thus, the proton motive force does not power gliding in *M. mobile*.

Arsenate ions compete with phosphate ions in substrate-level phosphorylation of ADP, and the resulting ADP-As molecule rapidly hydrolyzes (43). Therefore, we replaced the phosphate in our buffer with arsenate to study its effect. Arsenate caused a rapid and statistically significant drop in both the speed of gliding and the percentage of cells moving in the experiment (Fig. 3e and f). The drop in speed and fraction of cells moving is likely due to a specific effect of arsenate rather than simply the removal of phosphate since we were able to observe motility for extended periods of time in Tris- or TES [*N*-tris(hydroxymethyl)methyl-2-aminoethanesulfonic acid]-based buffers (data not shown). Presumably, the cells have a phosphate or ATP reserve that is rapidly abolished in the presence of arsenate. To further examine the effect of arsenate, we measured both ATP levels and gliding parameters from the onset of arsenate addition over an extended period of time. We found that both gliding speed and fraction of cells moving tracked the levels of ATP present in the cell (correlation coefficient,  $r^2 = 0.94$  and  $0.88$ , respectively) (Fig. 5). Thus, ATP

TABLE 1. Buffer formulations used in this study<sup>a</sup>

Test substance (concn in buffer)	Mode of action	Base buffer	Solvent (%)	Matched control buffer
Valinomycin (10 µM)	$K^+$ ionophore	Valinomycin buffer	MeOH (0.1)	Valinomycin buffer + 0.1% MeOH
FCCP (10 µM)	Protonophore	PBS/G	MeOH (0.1)	PBS/G + 0.1% MeOH
Arsenate (50 µM)	ATP poison	ArBS-K/G	Water	PBS-K/G
DCCD (10 µM)	$F_0F_1$ ATPase inhibitor	PBS/G	EtOH (0.025)	PBS/G + 0.025% EtOH
Sodium orthovanadate (1 mM)	P-type ATPase inhibitor	PBS-K/G	Water	PBS-K/G
Phenamil (10 µM)	Sodium channel blocker	PBS/G	MeOH (0.1)	PBS/G + 0.1% MeOH
Amiloride (10 µM)	Sodium channel blocker	PBS/G	Water	PBS/G
Rifampin (5 µg/ml)	RNA synthesis inhibitor	PBS/G	MeOH (0.1)	PBS/G + 0.1% MeOH
Puromycin (20 µM)	Protein translation inhibitor	PBS/G	Water	PBS/G

<sup>a</sup> Stock solutions were made of the test substance (for non-water-soluble compounds) in the solvent listed. The stock solution was then added to the base buffer, resulting in the final solvent concentration shown, where applicable. Abbreviations: MeOH, methanol; EtOH, ethanol.

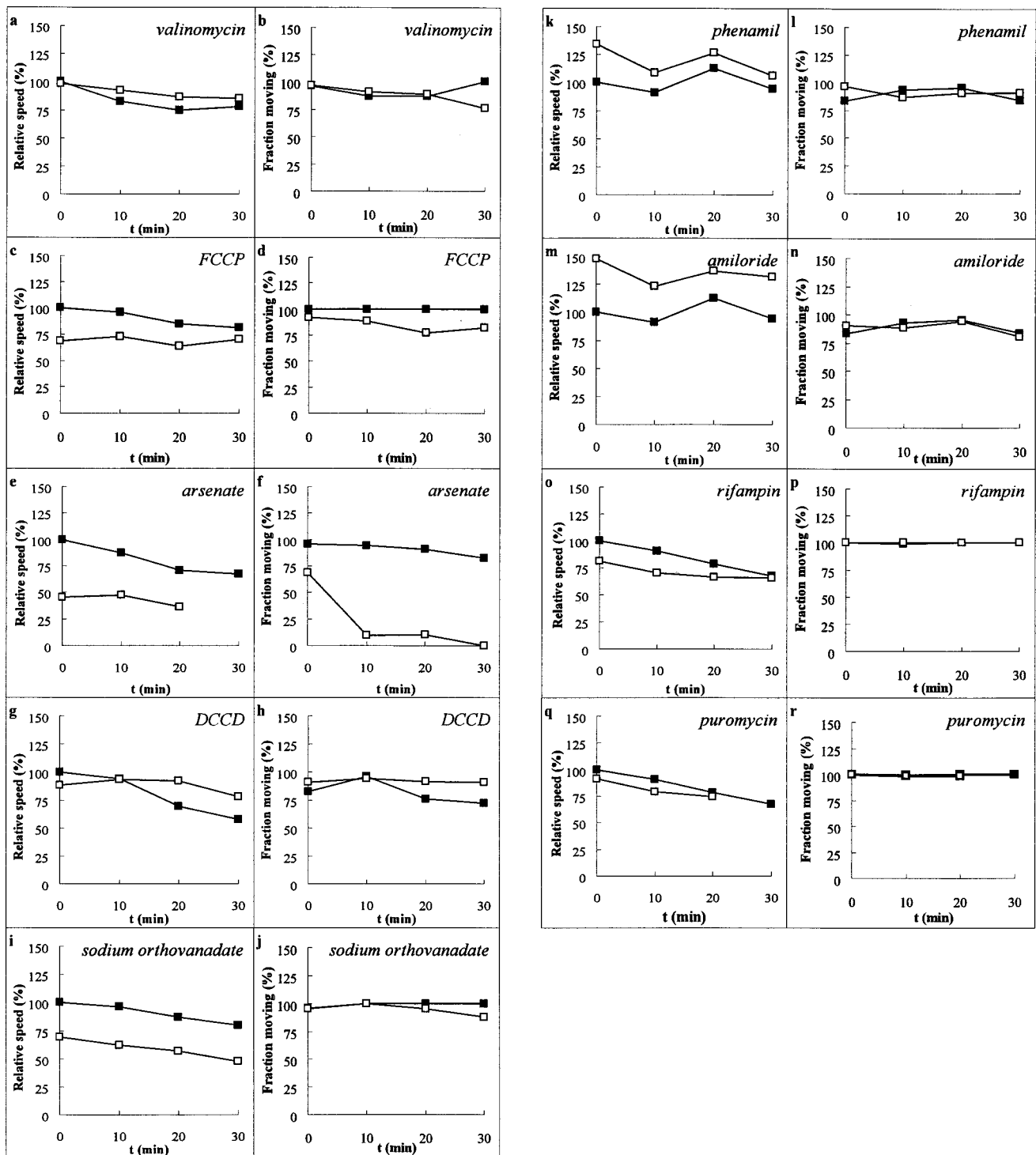


FIG. 3. Effect of ionophores, inhibitors, and poisons on gliding. Various agents were added (Table 1), and gliding was compared to that of a matched control. The graphs on the left-hand side of each pair show the speed of the cells that were moving at speeds greater than  $0.25 \mu\text{m/s}$  relative to the initial speed of the cells in the matched control. The graphs on the right-hand side of each pair show the fraction of cells moving. (a and b)  $10 \mu\text{M}$  valinomycin; (c and d)  $10 \mu\text{M}$  FCCP; (e and f)  $10 \text{mM}$  arsenate (note that no 30' test time point is shown [e] because no cells were moving at speeds greater than  $0.25 \mu\text{m/s}$ ); (g and h)  $10 \mu\text{M}$  DCCD; (i and j)  $1 \text{mM}$  sodium orthovanadate; (k and l)  $10 \mu\text{M}$  phenamil; (m and n)  $10 \mu\text{M}$  amiloride; (o and p)  $5 \mu\text{g}$  of rifampin/ml; (q and r)  $20 \mu\text{M}$  puromycin (note that the 30-min test time point was not taken due to technical difficulties). Filled squares are matched control preparations. Open squares are test preparations. Error bars are omitted for clarity, but standard deviations in speeds were generally  $\pm 25\%$ .

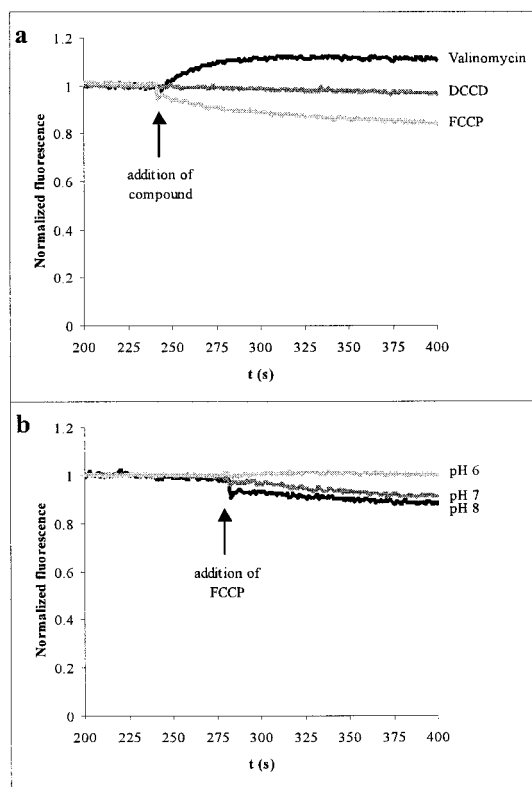


FIG. 4. Effect of selected substances on membrane potential. Cell suspensions were incubated with DiSC<sub>3</sub>[5] for 4 to 5 min, and then the indicated compound was added. Data are shown from 200 to 400 s of the experiment and are normalized to the average of data from the first 50 s of observations in the time period shown. Downward deflections indicate hyperpolarization. Upward deflections indicate depolarization. (a) Effect of 3  $\mu$ M valinomycin, 5  $\mu$ M FCCP, or 20  $\mu$ M DCCD in PBS-K/G (pH 8.0). The valinomycin preparation contained 50 mM KCl to match the gliding experiments. (b) Effect of FCCP at various pHs of PBS-K/G.

must be an important energy source for gliding motility in *M. mobile*.

Given the demonstrated importance of ATP for motility, we tested several ATPase inhibitors to see whether they might be involved in gliding motility. DCCD is an inhibitor of the F<sub>0</sub>F<sub>1</sub>-type proton translocating ATPase and is also reported to collapse  $\Delta p_H$  in some mycoplasmas (21, 22, 37). This compound had no effect on gliding motility (Fig. 3g and h) and lends further evidence that the proton motive force does not drive motility. The P-type Na<sup>+</sup> translocating ATPase inhibitor sodium orthovanadate caused a small but statistically significant decrease in gliding speed but did not reduce the percentage of cells gliding in a meaningful way (Fig. 3i and j) (39). To further examine the role of Na<sup>+</sup> gradients in gliding motility, we tested the sodium channel blockers phenamil and amiloride. Both of these compounds can abolish swimming motility in *Vibrio* species that utilize a sodium motive force to power flagellar rotation (2, 18). Neither of these compounds affected gliding motility in *M. mobile* (Fig. 3k to n), and it is therefore also unlikely that a sodium motive force ( $\Delta p_{Na}$ ) drives motility in *M. mobile* in the same manner that *Vibrio* species utilize  $\Delta p_{Na}$ .

Finally, we tested to see whether short-term biosynthesis of

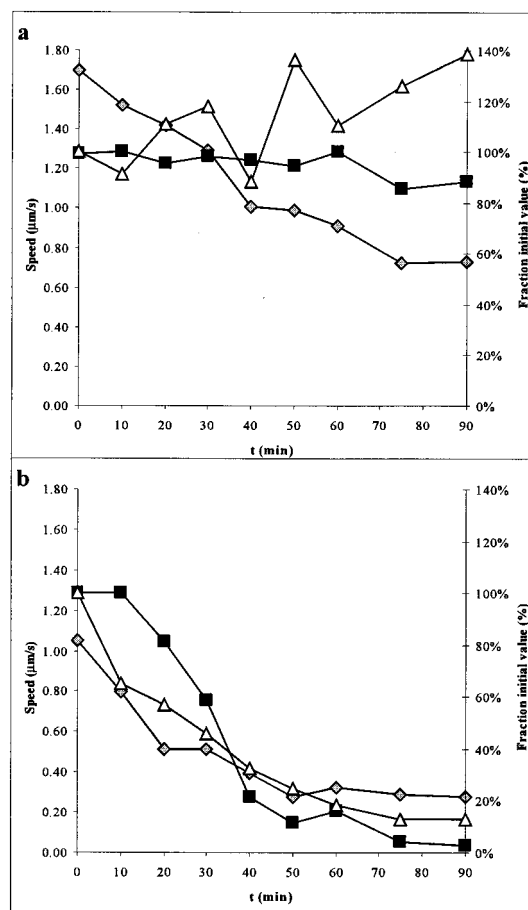


FIG. 5. Effect of arsenate on gliding motility and ATP levels. Parallel preparations were assayed for motility and ATP. (a) PBS-K/G (pH 8.0); (b) ArBS-K/G (50 mM arsenate). Filled squares indicate the fraction of cells moving (right axis), gray diamonds indicate gliding speed (left axis), and open triangles indicate the fraction of ATP remaining compared to the value at  $t = 0$  (right axis). Note that error bars have been omitted for clarity. Standard deviations in speeds were generally  $\pm 20\%$ , while standard deviations for ATP values were  $< 5\%$ .

protein was required for motility. We thought that this might be a reasonable possibility given that some microorganisms are reported to move by continuous extrusion of biomolecules or by continuous shedding of a protein trail during gliding motility (11, 40). We attempted to shut off transcription or translation in separate experiments. The transcriptional inhibitor rifampin prevents RNA synthesis, while the protein synthesis inhibitor puromycin mimics aminoacyl-tRNA and aborts translation (23). These compounds had no effect on motility over the 30-min span of the experiment (Fig. 3o to r).

### DISCUSSION

While there have been many remarkable observations of gliding motility of *M. mobile*, the source of energy for this phenomenon has not been convincingly demonstrated (15, 16, 28–30, 33–35). One study did examine the effects of various substances on *M. mobile* gliding (including some which overlap those discussed here), but that effort was directed at longer-term effects and did not specifically address the issue of ener-

getics (31). We studied the energy source by employing techniques and substances that were useful for investigating motility in many swimming bacteria. We hoped that elucidation of the energy source could guide further experimentation as to the mechanism of gliding motility.

We surmised that the energy for gliding motility in *M. mobile* might be supplied by an ion motive force (such as  $\Delta p_{\text{H}}$  or  $\Delta p_{\text{Na}}$ ) or that it could come directly from the hydrolysis of ATP. Many swimming bacteria (such as *E. coli*) utilize  $\Delta p_{\text{H}}$  as the source of power for motility, although some (such as *Vibrio alginolyticus*) use  $\Delta p_{\text{Na}}$  (24, 26). While mycoplasmas are certainly capable of and in fact do generate a proton motive force, it does not seem to supply the energy for gliding motility in this case. Rather, our observations suggest that motility is directly powered by the hydrolysis of ATP. Of the various compounds that we studied, arsenate was the only chemical that we found that had a dramatically significant effect on motility. We were able to successfully manipulate  $\Delta\Psi$  and  $\Delta\text{pH}$  of the cell but without effect on motility. However, we did observe a strong real-time correlation between ATP levels, gliding speed, and percentage of cells moving. We note that changes in ATP levels should eventually affect  $\Delta\Psi$  and  $\Delta\text{pH}$ ; however, this effect would occur on a time scale much longer than that for the changes induced by the addition of valinomycin or FCCP (compare Fig. 5b and 4a). We propose that the motor apparatus in *M. mobile*, which has yet to be identified, is most likely an ATPase or is directly coupled to an ATPase. This idea is strengthened by our observation that sodium orthovanadate caused a small but significant decrease in gliding speed. Vanadyl ( $\text{VO}_4^{3-}$ ) ions are well-known inhibitors of P-type  $\text{Na}^+/\text{K}^+$  translocating ATPases from a variety of organisms and are usually effective at concentrations 10- to 100-fold lower than those used here (6, 7). The ATPase motor system of *M. mobile* might be marginally sensitive to vanadate or might be derived from a P-type ATPase. This proposal may be helpful in targeting protein classes in *M. mobile* for further study with respect to gliding. Alternatively, it has been demonstrated that some sugar transport ABC-type ATPases are sensitive to vanadate, and the effect of vanadate in this experiment might be simply to limit the supply of fermentable sugar inside the cell (20).

*M. mobile* is also remarkable for its robust gliding motility compared to that in other mycoplasmas (14). Here, we show that motility in *M. mobile* is facilitated in part by protein components from the growth medium that condition glass surfaces to enable gliding. While our attempts to specifically identify the protein component by mass spectrometry were unsuccessful, we suspect that the essential protein(s) might act as a surfactant, allowing transient cytoadherence that would be necessary for gliding motility. Lipids might also be involved.

The mechanism of gliding motility in bacteria remains a mystery, and it is likely that several different schemes are employed in different genera. Variations on the basic mechanism might exist even within the closely grouped mycoplasmas. For instance, several large proteins implicated in surface adherence and motility in *M. mobile* have been identified (41). *Mycoplasma pneumoniae*, another gliding mycoplasma with a similar flask-shaped morphology, lacks any orthologs of these proteins. It is possible that there are different effectors of gliding motility even in these two closely related organisms.

Unfortunately, genetic manipulations have not yet been possible in *M. mobile* despite much success with mutagenesis techniques in other mycoplasmas (9, 10, 17, 42). Perhaps the upcoming report of the genome sequence of *M. mobile* combined with the power of comparative genomics will help to shed light on the mechanism(s) of gliding in the mycoplasmas (J. Jaffe et al., submitted for publication). Ultimately, it will be important to understand this new class of motor and its mechanism at the molecular level.

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