- Segall, J. E., S. M. Block, and H. C. Berg. 1986. Temporal comparisons in bacterial chemotaxis. *Proc. Natl. Acad. Sci. USA* 83:8987–8991.
- Shimizu, T. S., N. Le Novère, M. D. Levin, A. J. Beavil, B. J. Sutton, and D. Bray. 2000. Molecular model of a lattice of signalling proteins involved in bacterial chemotaxis. *Nature Cell Biol.* 2:1–5.
- Sourjik, V., and H. C. Berg. 2000. Localization of components of the chemotaxis machinery of *Escherichia coli* using fluorescent protein fusions. *Mol. Microbiol.* 37:740–751.
- Sourjik, V., and H. C. Berg. 2002a. Receptor sensitivity in bacterial chemotaxis. *Proc. Natl. Acad. Sci. USA* 99:123-127.
- Sourjik, V., and H. C. Berg. 2002b. Binding of the *Escherichia coli* response regulator CheY to its target measured in vivo by fluorescence resonance energy transfer. *Proc. Natl. Acad. Sci. USA* 99:12669–12674.

Rotary Motor

The structure of the rotary motor was described in Chapter 9 (Fig. 9.3) and its assembly was discussed in Chapter 10. Here, I will say more about function. Given that the diameter of the motor is less than one-tenth the wavelength of light and that it contains more than 20 of different kinds of parts (Appendix, Table A.3), it is a nanotechnologist's dream (or nightmare).

Power Source

dient; other cations and anions have been ruled out (Ravid and energy coupling, see Harold and Maloney (1996). $\Delta \psi \approx -120 \,\mathrm{mV}$, and $-59 \,\Delta pH \approx -50 \,\mathrm{mV}$. For cells grown at pH 7.7, Boltzmann's constant, T the absolute temperature, and e the the transmembrane pH difference (-2.3kT/e) ΔpH , where k is membrane electrical potential difference, $\Delta \psi$, and the other to force, Δp . In general, it comprises two terms, one due to the transin crossing the cytoplasmic membrane is called the protonmotive but rather by protons moving down an electrochemical graphate (ATP) the fuel that energizes muscles (Larsen et al., 1974), Flagellar motors of E. coli are not powered by adenosine triphos- $\Delta p \approx \Delta \psi \approx -140 \,\mathrm{mV}$. For a general discussion of chemiosmotic pH in the range 7.6 to 7.8. For cells grown at pH 7, $\Delta p \approx -170 \,\mathrm{mV}$ internal pH less the external pH. E. coli maintains its internal proton charge. At 24°C, 2.3 kT/e = 59 mV. By convention, $\Delta \psi$ is Eisenbach, 1984). The work per unit charge that a proton can do the internal potential less the external potential, and ΔpH is the

The dependence of speed on voltage has been measured in *E. coli* by wiring motors to an external voltage source. Filamentous cells were drawn roughly halfway into micropipettes, and the cytoplasmic membrane of the segment of the cell inside the pipette was made permeable to ions by exposure to the ionophore gram-

done with large markers (heavy loads) at speeds less than physiological range (up to -150 mV). These experiments were latter negative) caused the marker to spin (Fung and Berg, 1995). The rotation speed was directly proportional to Δp over the full recorded on videotape. Application of an electrical potential the segment of the cell outside the pipette, and its motion was icidin S. An inert marker was attached to a flagellar motor on rotation speed still appears proportional to Δp (Gabel and Berg markers (light loads) at speeds up to nearly 300Hz, and the between the external medium and the inside of the pipette (the 10 Hz. They have been repeated in a different way with small

organism can be starved and artificially energized, either with a and is sensitive to ionophores and uncouplers. Unlike E. coli, this gram-positive organism that lacks an endogenous energy reserve et al., 1975), a peritrichously flagellated, primarily fermentative, ing capacity, one can follow proton uptake by the increase in exterand shifting them to a medium with a lower concentration of potassium diffusion potential (by treating cells with valinomycin motors of the motile Streptococcus sp. strain V4051 (van der Drift cell vibration frequencies (Lowe et al., 1987). Finally, the data can can be determined by using power spectral analysis to measure nal pH. The frequency of rotation of filaments in flagellar bundles medium of lower pH). If this is done with a medium of low bufferpotassium ion) or with a pH gradient (by shifting cells to a tional to the initial swimming speed, as would be expected if a ping the motors by adding an antifilament antibody—this crossmotors. However, the two can be distinguished by suddenly stopinto the cell is much larger than the flux through its flagellar and the number of flagellar filaments per cell. The total proton flux be normalized to single motors by counting the number of cells uncertainty, due mainly to the difficulty of counting flagellar fixed number of protons carries a motor through each revolution the change in flux. This change was found to be directly proporlinks adjacent filaments in the fiagellar bundles—and measuring This number is about 1200 (Meister et al., 1987) but subject to The only measurement of proton flux that has been made is with

and Atsumi, 1989). Thus, flagellar motors are ion driven, not high pH, use sodium ions instead of protons (Imae, 1991; Imae Some bacteria, notably marine bacteria or bacteria that live at

> McCarter (2001) and Yorimitsu and Homma (2001). just proton driven. For reviews on sodium-driven motors, see

Torque-Generating Units

eight distinct proton channels (or pairs of proton channels), comduced eight equally spaced levels more than once, but never nine torque-generating units is that resurrections of this kind have proof rotation). The main argument for a complement of eight such torque (applies a similar force at the same distance from the axis unit (comprising MotA and MotB) adds the same increment of cells increases in a number of equally spaced steps, as shown in plasmids. As new protein is synthesized, the speed of tethered this was obtained by restoring the motility of paralyzed cells prising one or more copies of the proteins MotA and MotB Fig. 12.1. This indicates that each additional torque-generating (mot mutants) via the expression of wild-type genes carried by (currently thought to be four MotA and two MotB). Evidence for The flux through the flagellar motor is divided into as many as

Stepping

ments tend to untwist, carrying the cell body forward. Therefore, elastic elements, comprising the rod, hook, and filament) causes structure linking the rotor to the tethering surface (a series of occur with exponentially distributed waiting times. We have been proton passage is likely to occur at random times, the steps will other words, the motor should behave like a stepping motor. Since by a fixed increment, once the tether relaxes (see below). In moves a torque generator (a MotA, MotB complex) one step (one It is likely that the passage of each proton (or each proton pair) smoothed out. To succeed, one probably needs to work at reduced discontinuities in the relative motion of rotor and stator are that structure to twist. When less torque is applied, these ele-The main reason, advanced then, is that the torque applied to the looking for such steps since 1976 (Berg, 1976) but without success. the cell wall. As this linkage relaxes, a tethered cell should rotate the components that link that generator to the rigid framework of binding site) along the periphery of the rotor, suddenly stretching

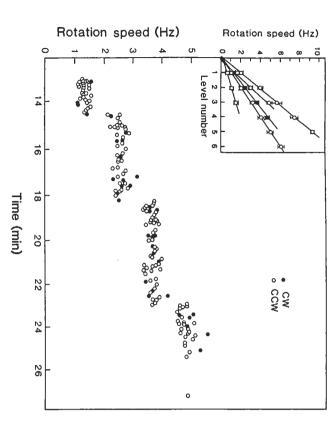


FIGURE 12.1. Rotation speed of a tethered motA cell, E. coli strain MS5037(pDFB36), following addition (at time 0) of the inducer isopropyl β-p-thiogalactoside (IPTG, added in a minimal medium containing glycerol, glucose, and essential amino acids). Filled circles indicate CW rotation, open circles CCW rotation. The inset shows the mean rotation speed (±standard error of the mean) at each level (step of the staircase) as a function of level number, for this cell (closed circles) and for four additional cells (open circles). (Blair & Berg, 1988, Fig. 1, reprinted with permission from the American Association for the Advancement of Science.)

torque, for example, with a one-generator motor driving a small viscous load, perhaps just a hook. Such an object is expected to spin quite rapidly, so the technical problems are formidable.

One route around this difficulty is to examine variations in rotation period. If n steps occur at random each revolution, then the ratio of the standard deviation to the mean should be $n^{-1/2}$ (see the appendix in Samuel and Berg, 1995), so one can determine n. With tethered wild-type cells, the answer turns out to be about 400. This work also showed that tethered cells are not free to execute rota-

tional brownian motion. Thus, the rotor and stator are interconnected most of the time.

others, but that each remains connected to the rotor most of the number is 100 per revolution. If m = 8, the apparent step number rotor is half of what it would be at m = 1, so the apparent step not only that each force generator steps independently of all the apparent step number would remain 50. So this experiment argues is detached most of the time (for most of its duty cycle), then the is 400 per revolution. If, on the other hand, each torque generator ators that have not stepped. If m = 2, the net movement of the rotor, it also must stretch the linkages of the m-1 torque generof the cell wall, then when that linkage relaxes and moves the one steps, suddenly stretching its linkage to the rigid framework not just 50? If m torque generators are attached to the rotor and the time (see below). time. In fact, the torque generators must be attached nearly all of going resurrection (as in Fig. 12.1), and the number of steps per the rotor, say 50, then why is the number of steps per revolution torque generators interact with a fixed number of binding sites on increasing by about 50 steps per level (Samuel and Berg, 1996). If revolution was found to increase linearly with level number This stochastic analysis was repeated with tethered cells under-

Torque-Speed Dependence

A crucial test of any motor model is its torque-speed dependence. Measurements of the torque generated by motors of *E. coli* have been made over a wide range of speeds, including speeds in which the motor is driven backward, with the results shown in Fig. 12.2 (thick lines). At 23°C, the torque exerted by the motor is approximately constant, all the way from negative speeds of at least –100Hz to positive speeds of nearly 200Hz. At higher speeds it declines approximately linearly, crossing the 0-torque line at about 300Hz. At lower temperatures, the region of transition from constant torque to declining torque—we call this the "knee"—shifts to lower speeds, and the region of decline steepens (Berg and Turner, 1993; Chen and Berg, 2000a); the latter parts of the curves can be mapped onto one another with scaling of the speed axis.

Estimates of the torque generated in the low-speed regime range from about 2.7×10^{-11} dyn cm (2700 pN nm) to 4.6×10^{-11} dyn cm (4600 pN nm), the smaller value from estimates of the viscous

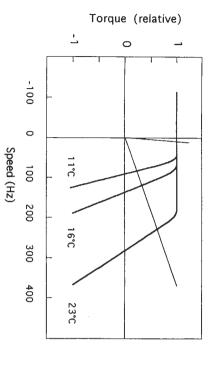


FIGURE 12.2. The torque-speed curve for the flagellar motor of *E. coli* shown at three temperatures (thick lines), together with two load lines (thin lines), one for an object the size of the cell body of wild-type *E. coli* (effective radius about 1 µm, left), the other for a latex bead of radius about 0.3 µm, right. (Adapted from Fig. 16 of Berg and Turner, 1993.) Later work showed that the torque declines somewhat in the low-speed regime, by about 10% between stall and the knee; see the text.

drag on tethered cells of *Streptococcus* (Lowe et al., 1987), and the larger value from the force exerted by tethered cells of *E. coli* on latex beads held in an optical trap (Berry and Berg, 1997).

a geometrical factor times the bulk viscosity, η , where η is independent of Ω (independent of the rate of shear). For an isolated vinylpyrrolidone (Berg and Turner, 1979). For such a medium, f is long unbranched molecules, such as methylcellulose or polydition satisfied in a dilute aqueous medium that does not contain slope f. Here, we assume that the medium is Newtonian, that is plot, this function is a straight line passing through the origin, with ity, Ω (2π times its rotation speed, in Hz). In a torque versus speed required to rotate an object of fixed shape in a viscous medium is at the right for a small object. To see this, note that the torque, N will spin at the speed at which the torque generated by the motor that the frictional drag coefficient does not depend on Ω , a conits rotational frictional drag coefficient, f, times its angular veloc-12.2 (thin lines), the one at the left for a large object and the one is balanced by the torque exerted on the object by viscous drag This torque is defined by load lines, such as those shown in Fig. A motor driving an inert object (a cell body, a latex bead, etc.)

sphere of radius a spinning about an axis through its center, for example, this geometrical factor is $8\pi a^3$. For compact globular objects, the actual shape is not very critical; however, accurate values can be computed (Garcia de la Torre and Bloomfield, 1981). The distance from the tethering surface does not really matter, either, provided that the gap between the object and the surface is at least 0.2 cell radii (Berg, 1976; Jeffery, 1915).

At 23°C and for the load line shown at the left in Fig. 12.2, the motor runs at 10Hz; for the load line shown at the right, it runs about 220Hz. For a very shallow load line (e.g., one for a free hook), the speed would be close to the zero-torque speed, about 290Hz. A motor free-running in this way always operates in the upper-right-hand quadrant of Fig. 12.2. It cannot drive itself backward, although it can redefine what is meant by forward by switching from counterclockwise (CCW) to clockwise (CW) or back again. Nor can it spin faster than its speed at zero load. To probe the upper-left-hand or lower-right-hand quadrants of Fig. 12.2, one needs to subject the motor to torque applied externally.

externally applied torque before the break occurred. That differwhich the cell body turns at a given value of externally applied the C-ring is sheared off of the bottom of the rotor (Fig. 9.3), drops to zero, the cell appears free to execute rotational brownto spin CW) often breaks catastrophically: motor torque suddenly backward (CW if it is trying to spin CCW, or CCW if it is trying For reasons that we do not understand, the motor of a cell driven or backward. Speeds of several hundred Hz are readily attainable field. Therefore, it is possible to spin a tethered cell either forward and changes sign with changes in the direction of rotation of that torque varies as the square of the magnitude of the electric field applied torque is the cross-product of these vectors. The applied ization vector leads or lags the electric-field vector. The externally the finite time required for redistribution of charges, the polarthe same direction as the applied electric field. However, due to dipole field due to the polarization rotates at the same rate and in latter reference, the external electric field polarizes the cell. The rotating electric field (Berg and Turner, 1993). As explained in the Cells were tethered and exposed to a high frequency (2.25MHz) torque with the speed at which it turned at the same value of intact. Once the motor has broken, one can compare the speed at disengaging all torque-generating units but leaving the bearings ian motion, and the motor fails to recover. Our best guess is that One way to do this is by electrorotation (Washizu et al., 1993).

speed at which it turned when intact. The data shown by the thick ence is proportional to the torque generated by the motor at the lines in Fig. 12.2 were determined in this way.

or translocation of protons are not limiting. In the high-speed gized cells of Streptococcus (Khan and Berg, 1983). Evidently, at effects were relatively small, as found earlier for artificially enersizes were attached to the flagellar stubs, and the slopes of their adsorbed onto positively charged glass. Latex beads of various high speed is due to limits in rates of proton transfer (protor 2000b). This is what one would expect if the decline in torque at Fig. 12.2, and solvent isotope effects were large (Chen and Berg regime, torque was strongly temperature dependent, as seen in where rates of displacement of internal mechanical components low speeds, the motor operates near thermodynamic equilibrium torque was independent of temperature, and solvent isotope found to drop by about 10% from stall to the knee. In this regime load lines were increased by addition of the viscous agent Ficol lines. Flagella were shortened by viscous shear, and cells were right-hand quadrant of Fig. 12.2 was done by manipulating load (Chen and Berg, 2000a). In the low-speed regime, torque was Additional work on the behavior of the motor in the upper-

narrow, and why the position of that boundary is so sensitive to boundary between the low-speed and high-speed regimes is so understand why the low-speed regime is so broad, why the torque (Berry and Berg, 1999). Eventually, we would like to step that is effectively irreversible and insensitive to external rotation rules out models (e.g., thermal ratchets) that contain a erstroke mechanism, while the absence of a barrier to backward and dissipates most of the available free energy, that is, for a powmodel in which the rate-limiting step depends strongly on torque temperature. Slowly declining torque in the low-speed regime argues for a

revolution and speed 10 Hz, the proton flux is $1.2 \times 10^4 \,\mathrm{s}^{-1}$. For charge times protonmotive force. Assuming 1200 protons per rate at which protons can do work, is proton flux times proton and speed 10 Hz, this is $2.9 \times 10^5 \,\mathrm{pN}\,\mathrm{nm}\,\mathrm{s}^{-1}$. The power input, the tains rotation at angular velocity Ω , is N Ω . For torque 4600 pN nm $(1.2 \times 10^4 \text{s}^{-1})$ (e) $(0.17 \text{ V}) = 2.0 \times 10^3 \text{ eV s}^{-1}$. Since 1 eV (one elec E. coli at pH 7, $\Delta p \approx -170 \,\mathrm{mV}$. Therefore, the power input is The power output, the power dissipated when a torque N sus

> motor, power output divided by power input, is about 90%. Within been measured in E. coli---the efficiency could be 1. 10⁵ pN nm s⁻¹. So, by this crude estimate, the efficiency of the tron volt) = 1.6×10^{-12} erg = 160 pN nm, the power input is 3.2×10^{-12} the uncertainty of the measurements—the proton flux has not

stant up to the knee, and then it declines. There is no discontinuwith speed. Therefore, the efficiency remains approximately conthrough each revolution, the power input also increases linearly then it declines. If a fixed number of protons carries the motor boundary between the low-speed and high-speed regimes, and the F₀-ATPase pumps protons when driven backward by F₁. 1997). As the motor turns backward, it must pump protons, just as ity in torque as one crosses the zero-speed axis (Berry and Berg The power output, $N\Omega$, increases linearly with speed up to the

and lipid, about 2). So almost any kind of chemistry will do. in a medium of dielectric constant 40 (midway between water, 80 in magnitude to that between two electrons 4.8 angstroms apart generating units, then each contributes 25 pN. This is a force equathen 200 pN is applied. If there are eight independent forceunits act at the periphery of the rotor at a radius of about 20nm the stall torque of 4000 pN nm and assume that force-generating but not large on an absolute scale. If we take a ballpark figure for The force exerted by each force-generating unit is substantial

as large, and $F \approx 22 \,\mathrm{pN}$. So, given the estimate of 25 pN per force two protons are required per elementary step, the force is twice subunits) and a rotor radius of 20 nm, d \approx 2.4 nm. So $F \approx 11$ pN. If Assuming 52 steps per revolution (twice the number of FliG d is the displacement generated by the transit of one proton force-generator can do, Fd, where F is the force that it exerts, and or 27 pN nm. At unit efficiency, this equals the work that the trochemical gradient is $e\Delta p$. Given $\Delta p \approx -170 \,\mathrm{mV}$, this is $0.17 \,\mathrm{eV}$ step is likely. generating unit made above, the displacement of two protons per The energy available from one proton moving down the elec-

Angular Dependence of Torque

or to allow them to turn slowly forward (Berry & Berg, 1997), torque did not vary appreciably with angle. A very different resul When optical tweezers were used to drive cells slowly backward

is obtained when one energizes and de-energizes tethered cells and asks where they stop or watches them spin when the proton-motive force is very low. When this was done with *Streptococcus*, periodicities were observed of order 5 or 6 (Khan et al., 1985). This probably reflects small periodic barriers to rotation intrinsic to the bearings.

Duty Ratio

angle. I hus, the torque-generating unit would not be able to keep rad s⁻¹, and $\alpha = 2.5 \times 10^4$ s⁻¹. So, in 1.6×10^{-5} s, the twist in the tether as a sphere of radius $a = 20 \,\mathrm{nm}$ immersed in a medium of viscos rotational frictional drag coefficient. If we approximate the rotor initial twist, and α is the torsional spring constant divided by the duty ratio is 0.999, so that the torque-generating unit detaches for spinning ~1.2 Hz (Fig. 12.1), the step interval is 1.6×10^{-2} s. If the so if the torque-generating unit lets go, the tether will unwind 38 degrees, or by 19 degrees, that is, by more than twice the step decreases from 57 degrees to 57 $\exp(-2.5 \times 10^4 \text{ s}^{-1} \times 1.6 \times 10^{-5} \text{ s}) =$ then the frictional drag coefficient, $8\pi\eta a^3$, is 2×10^{-16} dyn cm per ity $\eta = 1 P (1 \text{g cm}^{-1} \text{s}^{-1})$, which is about right for a lipid membrane revolution, the displacement is 7.2 degrees per step. If the cell is driving the rotor backward. If the single unit steps 50 times per drag on the cell body is enormous compared to that on the rotor. tether is twisted up about 1 radian, or 57 degrees. Now the viscous hook—is about 5×10^{-12} dyn cm rad⁻¹ (Block et al., 1989), so the spring constant of the tether-most of the compliance is in the motor generates a torque of about 5×10^{-12} dyn cm. The torsional torque of about 4×10^{-11} dyn cm (4000 pN nm), then the single-unit wild-type motor with eight torque-generating units generates a unit, as in the first step of the resurrection shown in Fig. 12.1. If a sider a tethered cell being driven by a single torque-generating torque generators, like molecules of kinesin, are processive. Conargument shows that the duty ratio must be close to 1. Evidently, number of torque generators, as observed, if each torque genera- 1.6×10^{-5} s during each cycle, how far will the tether unwind? the torque-generating units had a high duty ratio. The following tor remained attached to the rotor most of the time, that is, if apparent number of steps per revolution would increase with the The tether unwinds exponentially: $\theta = \theta_0 \exp(-\alpha t)$, where θ_0 is the In our stochastic analysis of steps (above) we argued that the

up. So the duty ratio must be close to 1. The interaction between the torque-generating unit and the rotor must be such that the rotor is not able to slip backward. If one imagines that a torque-generating unit binds to successive sites along the periphery of the rotor, then it has no unbound states. If each torque-generating unit has two proton channels (Braun and Blair, 2001), it is possible that a MotA associated with one channel remains attached to a FliG, while the MotA associated with the other channel takes the next step.

Switching

Finally, the motor can run in either direction with approximately equal efficiency. Although the force-generating elements move independently, they all switch at the same time: changes in direction occur in an all-or-none fashion within a few milliseconds. Evidently, the rotor suddenly changes shape, so that the force-generating elements step along a different track. What sort of change in conformation occurs? And why is this process so sensitive to the concentration of CheY-P?

Models

The fundamental question is how the flagellar motor generates torque, namely, how inward motion of one or more ions through a torque-generating unit causes it to advance circumferentially along the periphery of the rotor. Once that is understood, the nature of the conformational change required for switching, namely, how the direction of advance is distinguished from that of retreat, is likely to be self-evident.

Moving parts of the motor are submicroscopic and immersed in a viscous medium (water or lipid), so the Reynolds number is very small (see Chapter 6). And everything is overdamped (Howard, 2001, pp. 41–45). Thus, the designer does not have the benefit of flywheels or tuning forks. If, for example, the operator of the motor driving a tethered cell of *E. coli* 10 Hz were to put in the clutch, the cell body would coast no more than a millionth of a revolution. So if there is a stage in the rotational cycle in which the torque changes sign, the motor will stop. Predicting net torque after averaging over a complete cycle is not sufficient. And mech-

anisms in which energy is stored in vibrational modes are not viable. However, one can use energy available from an electrochemical potential to stretch a spring and then use that spring to apply a steady force. As we have seen, the force required is modest, and almost any kind of chemistry will do.

a fixed number of protons carry the motor through each revoluexperiment has attempted to measure this flux (Meister et al., protons traversing the motor equal to the mechanical work that it operate reversibly at unit efficiency, with the free energy lost by namic equilibrium. For slow rotation near stall, the motor must energy is dissipated; therefore, the stalled motor is at thermodytion. The running torque at low speeds is close to the stall torque protons flow through the motor when it is stalled, this implies that of the rotor is driven by a proton (or sodium-ion) flux. Only one which the motor is tightly coupled. range, as observed. So the evidence is consistent with a model in portional to the protonmotive force over its full physiological performs. This implies that the torque near stall should be pro-1987), and flux and speed were found to be linearly related. Unless (Fig. 12.2). If the motor is stalled and no protons flow, no free Motion of the torque-generating units relative to the periphery

cell by backward rotation, and steep barriers are not expected. In generating torque (e.g., by enabling a ratchet that is powered theravailable from the electrochemical gradient and rotational work electrochemical gradient is directly involved in generating torque, stroke mechanism. expected. So the torque-speed curves of Fig. 12.2 favor a powerward torques are applied, and barriers to backward rotation are gradient is small, so the system must wait, even when large backpling, the likelihood of transit of ions against the electrochemical erate large changes in speed. In the ratchet case, with tight courelatively flat plateau, because small changes in torque can genmally). In the powerstroke case, protons can be driven out of the occur synchronously, or whether the ion is indirectly involved in that is, participates in a powerstroke in which dissipation of energy then the torque-speed curve (as plotted in Fig. 12.2) can have a addition, if the rate-limiting step is strongly torque dependent, An important question is whether the ion that moves down the

There appear to be essential electrostatic interactions between specific residues in the cytoplasmic domain of MotA and the C-terminal domain of FliG (Zhou et al., 1998a). Here, charge

complementarity is more important than surface complementarity; that is, long-range interactions appear to be more important than tight binding. Since some models for torque generation require transfer of protons from the stator to the rotor, it was expected that acidic residues on FliG might be more important than basic residues. However, replacement of the acidic residues deemed important for torque generation with alanine still allowed some rotation, while reversing their charge had a more severe effect (Lloyd and Blair, 1997). An extension of this study failed to identify any conserved basic residues critical for rotation in MotA, MotB, FliG, FliM, or FliN, and only one conserved acidic residue critical for rotation, Asp32 of MotB (Zhou et al., 1998b). Other alternatives were considered and either ruled out or deemed unlikely. Therefore, the only strong candidate for a residue that functions directly in proton conduction is Asp32 of MotB.

MotA and MotB appear to form a cassette containing a transmembrane channel that supports proton flow, generating transformations that drive movement along the periphery of the rotor. That the ion-dependence is determined solely by MotA and MotB (or their homologs) has been shown conclusively in recent experiments in which transmembrane and cytoplasmic domains of MotA and MotB were replaced by homologous parts of PomA and PomB, from *Vibrio alginolyticus*. With only the C-terminal periplasmic domain of MotB remaining, the *E. coli* motor became sodium-ion driven rather than proton driven (Asai et al., 2003).

Given the above work, I would bet on a cross-bridge mechanism of the kind that Blair and colleagues propose (Braun et al., 1999; Kojima and Blair, 2001). In such a scheme, proton transport drives a cyclic sequence in which (1) a proton binds to an outward-facing binding site; (2) the protonmotive force drives a conformational change, a powerstroke that moves the rotor forward (or stretches a spring that moves it forward) and transforms the binding site to an inward-facing site; and (3) proton dissociation triggers detachment of the cross-bridge from the rotor, its relaxation to the original shape, and reattachment to an adjacent site. If the MotA/MotB complex is two-headed, one head could remain attached while the other stepped, thus ensuring a high duty ratio.

But to be honest, we really do not understand how the motor works, i.e., how proton translocation generates torque. Modeling would help, but what is needed most is more structural information.

and Blair (2003). For a catalog of early models, see Berg and material in this chapter was adapted from Berg (2003). motor models, see Berry (2000) and Bustamante et al. (2001). The Turner (1993). For tutorials on the mathematical treatment of Berry and Armitage (1999), Berry (2000, 2003), Berg (2000, 2003), motors, see Läuger and Kleutsch (1990), Caplan and Kara-Ivanov (1993), Schuster and Khan (1994), Macnab (1996), Khan (1997). For other reviews on the structure and function of proton-driven

References

- Asai, Y., T. Yakushi, I. Kawagishi, and M. Homma. 2003. Ion-coupling determinants of Na⁺-driven and H⁺-driven flagellar motors. J. Mol Biol. 327:453-463.
- Berg, H. C. 1976. Does the flagellar rotary motor step? In: Cell Motility, T. Pollard, J. Rosenbaum, editors. Cold Spring Harbor Laboratory, Cold Spring Harbor Conferences on Cell Proliferation. R. Goldman, Cold Spring Harbor, NY, pp. 47-56.
- Berg, H. C. 2000. Constraints on models for the flagellar rotary motor Philos. Trans. R. Soc. Lond. B 355:491-501.
- Berg, H. C. 2003. The rotary motor of bacterial flagella. Annu. Rev. Biochem. 72:19-54.
- Berg, H. C., and L. Turner. 1979. Movement of microorganisms in viscous environments. Nature 278:349-351.
- Berg, H. C., and L. Turner. 1993. Torque generated by the flagellar motor of Escherichia coli. Biophys. J. 65:2201-2216.
- Berry, R. B. 2000. Theories of rotary motors. Philos. Trans. R. Soc. Lond B 355:503-509.
- Berry, R. B. 2003. The bacterial flagellar motor. In: Molecular Motors. M. Schliwa, editor. Wiley-VCH, Weinheim, pp. 111-140.
- Berry, R. B., and J. P. Armitage. 1999. The bacterial flagella motor. Adv Microbiol. Physiol. 41:291-337.
- Berry, R. M., and H. C. Berg. 1997. Absence of a barrier to backwards rotation of the bacterial flagellar motor demonstrated with optical tweezers. Proc. Natl. Acad. Sci. USA 94:14433-14437.
- Berry, R. M., and H. C. Berg. 1999. Torque generated by the flagellar motor of Escherichia coli while driven backward. Biophys. J. 76:
- Blair, D. F. 2003. Flagellar movement driven by proton translocation
- Blair, D. F., and H. C. Berg. 1988. Restoration of torque in defective fla gellar motors. Science 242:1678-1681.

- Block, S. M., D. F. Blair, and H. C. Berg. 1989. Compliance of bacterial flagella measured with optical tweezers. Nature 338:514-517.
- Braun, T. F., and Blair, D. F. 2001. Targeted disulfide cross-linking of the stator complex. Biochemistry 40:13051-13059. MotB protein of Escherichia coli: evidence for two H+ channels in the
- Braun, T. F., S. Poulson, J. B. Gully, et al. 1999. Function of proline residues of MotA in torque generation by the flagellar motor of *Escherichia coli. J. Bacteriol.* 181:3542–3551.
- Bustamante, C., D. Keller, and G. Oster. 2001. The physics of molecular motors. Acc. Chem. Res. 34:412-420.
- Caplan, S. R., and M. Kara-Ivanov. 1993. The bacterial flagellar motor Int. Rev. Cytol. 147:97-164.
- Chen, X., and H. C. Berg. 2000a. Torque-speed relationship of the flagellar rotary motor of Escherichia coli. Biophys. J. 1041. 78:1036-
- Chen, X., and H. C. Berg. 2000b. Solvent-isotope and pH effects on flagellar rotation in Escherichia coli. Biophys. J. 78:2280-2284.
- Fung, D. C., and H. C. Berg. 1995. Powering the flagellar motor of Escherichia coli with an external voltage source. Nature 375:809-812.
- Gabel, C. V., and H. C. Berg. 2003. The speed of the flagellar rotary motor of Escherichia coli varies linearly with protonmotive force. Proc. Natl Acad. Sci. USA 100:8748-8751.
- Garcia de la Torre, J., and V. A. Bloomfield. 1981. Hydrodynamic propcations. Q. Rev. Biophys. 14:81-139. erties of complex, rigid, biological macromolecules: theory and appli-
- Harold, F. M., and P. C. Maloney. 1996. Energy transduction by ion cur-Press, Washington DC, pp. 283-306. Biology. F. C. Neidhardt, R. Curtiss, J. L. Ingraham, et al., editors. ASM rents. In: Escherichia coli and Salmonella: Cellular and Molecular
- Howard, J. 2001. Mechanics of Motor Proteins and the Cytoskeleton Sinaur Associates, Sunderland, MA.
- Imae, Y. 1991. Use of Na+ as an alternative to H+ in energy transduction. Tokyo, pp. 197-221. In: New Era of Bioenergetics. Y. Mukohata, editor. Academic Press
- Imae, Y., and T. Atsumi. 1989. Na⁺-driven bacterial flagellar motors. J. Bioenerg. Biomembr. 21:705-716.
- Jeffery, G. B. 1915. On the steady rotation of a solid of revolution in viscous fluid. Proc. Lond. Math. Soc. 14:327-338.
- Khan, S. 1997. Rotary chemiosmotic machines. Biochim. Biophys. Acta 1322:86–105.
- Khan, S., and H. C. Berg. 1983. Isotope and thermal effects in chemiosmotic coupling to the flagellar motor of Streptococcus. Cell 32:913-
- Khan, S., M. Meister, and H. C. Berg. 1985. Constraints on flagellar rotation. J. Mol. Biol. 184:645-656.

Larsen, S. H., J. Adler, J. J. Gargus, and R. W. Hogg. 1974. Chemome-chanical coupling without ATP: the source of energy for motility and chemotaxis in bacteria. *Proc. Natl. Acad. Sci. USA* 71:1239–1243.

Läuger, P., and B. Kleutsch. 1990. Microscopic models of the bacterial flagellar motor. *Comments Theor. Biol.* 2:99–123.

Lloyd, S. A., and D. F. Blair. 1997. Charged residues of the rotor protein FliG essential for torque generation in the flagellar motor of *Escherichia coli. J. Mol. Biol.* 266:733-744.

Lowe, G., M. Meister, and H. C. Berg. 1987. Rapid rotation of flagellar bundles in swimming bacteria. *Nature* 325:637–640.

Macnab, R. M. 1996. Flagella and motility. In: *Escherichia coli* and *Salmonella*: Cellular and Molecular Biology. F. C. Neidhardt, R. Curtiss, J. L. Ingraham, et al., editors. ASM Press, Washington, DC, pp. 123–145.

McCarter, L. L. 2001. Polar flagellar motility of the Vibrionaceae. Microbiol. Mol. Biol. Rev. 65:445-462.

Meister, M., G. Lowe, and H. C. Berg. 1987. The proton flux through the bacterial flagellar motor. *Cell* 49:643–650.

Ravid, S., and M. Eisenbach. 1984. Minimal requirements for rotation of bacterial flagella. *J. Bacteriol.* 158:1208–1210.

Samuel, A. D. T., and H. C. Berg. 1995. Fluctuation analysis of rotational speeds of the bacterial flagellar motor. *Proc. Natl. Acad. Sci. USA* 92: 3502–3506.

Samuel, A. D. T., and H. C. Berg. 1996. Torque-generating units of the bacterial flagellar motor step independently. *Biophys. J.* 71:918–923.

Schuster, S. C., and S. Khan. 1994. The bacterial flagellar motor. Annu Rev. Biophys. Biomol. Struct. 23:509-539.

van der Drift, C., J. Duiverman, H. Bexkens, and A. Krijnen. 1975. Chemotaxis of a motile *Streptococcus* toward sugars and amino acids. *J. Bacteriol.* 124:1142–1147.

Washizu, M., Y. Kurahashi, H. Iochi, et al. 1993. Dielectrophoretic measurement of bacterial motor characteristics. *IEEE Trans. Ind.* Appl. 29:286–294.

Yorimitsu, T., and M. Homma. 2001. Na⁺-driven flagellar motor of *Vibrio*. *Biochim. Biophys. Acta* 1505:82–93.

Zhou, J., S. A. Lloyd, and D. F. Blair. 1998a. Electrostatic interactions between rotor and stator in the bacterial flagellar motor. *Proc. Natl. Acad. Sci. USA* 95:6436–6441.

Zhou, J., L. L. Sharp, H. L. Tang, et al. 1998b. Function of protonatable residues in the flagellar motor of *Escherichia coli*: a critical role for Asp 32 of MotB. *J. Bacteriol.* 180:2729–2735.

Epilogue

What We Have Learned

I have told you some things about a free-living organism only one micron in size. It is equipped with sensors that count molecules of interest in its environment, coupled to a readout device that computes whether these counts are going up or down. The output is an intracellular signal that modulates the direction of rotation of a set of rotary engines, each turning a propeller with variable pitch. Each engine (or motor) is driven, in turn, by several forcegenerating elements (like pistons), powered by a transmembrane ion flux. In addition to a gear shift (labeled forward and reverse but prone to shift on its own) there is a stator, a rotor, a drive shaft, a bushing, and a universal joint.

We know a great deal about what all this machinery does for the bacterium, a fair amount about the structures of the molecular components involved (particularly those that have been crystallized), and even how the organism programs their syntheses. We know less about the precise ways in which these components function.

Levels of Amazement

Some wonder how the flagellar motor possibly could have evolved. The problem here is that we do not know about earlier states. What was the flagellar motor doing, for example, before the acquisition of the propeller (if, indeed, that was the sequence of events)? Perhaps it was winding up DNA. Or maybe it was injecting toxins into other cells as part of a program of conquest. In any event, it must have been doing something that promoted the survival of the organism. Evolution is opportunistic: it builds on components already at hand. One can not turn off the organism