

Macro-Level and Genetic-Level Responses of *Bacillus subtilis* to Shear Stress

Susmita Sahoo,[†] Rajesh K. Verma,[†] A. K. Suresh,^{†,‡} K. Krishnamurthy Rao,[‡] Jayesh Bellare,^{†,‡} and G. K. Suraishkumar^{*,†,‡}

Department of Chemical Engineering and School of Biosciences and Bioengineering, Indian Institute of Technology, Bombay, Powai, Mumbai 400076 India

Responses of bacterial (*Bacillus subtilis*) cells under different shear levels, from both the macro and genetic viewpoints, have been presented. The responses were studied using a novel, couette flow bioreactor (CFB), in which the entire cultivation can be performed under defined shear conditions. Oxygen supply, the normal limiting factor for entire cultivations under defined shear conditions, has been achieved by passing air through a poly(tetrafluoroethylene) (PTFE) membrane fixed on the inner cylinder of the CFB. More importantly, analyses of the oxygen transfer capabilities as well as the shear rates show that in this CFB, the effects of defined shear can be studied without interference from the effects of oxygen supply. Further, the shake flask can be used as a proper control for studying the shear effects, mainly because the shear rate in the shake flask under normal shaker operating conditions of 190 rpm has been estimated to be a negligible 0.028 s^{-1} compared to a value of 445 s^{-1} at the lowest rpm employed in the CFB. At the macro level the cell size decreased by almost 50% at 1482 s^{-1} compared to that at 0.028 s^{-1} , the growth rate increased by 245%, and the maximum cell concentration increased by 190% when the shear rate was increased from 0.028 to 1482 s^{-1} . The specific intracellular catalase level increased by 335% and protease by 87% at 1482 s^{-1} as compared to the control cultures at a shear rate 0.028 s^{-1} . In addition, the specific intracellular reactive oxygen species level (siROS) at the highest shear rate was 9.3-fold compared to the control conditions. At the genetic level we have established the involvement of the transcription factor, σ^B , in the bacterial responses to shear stress, which was unknown in the literature thus far; the σ^B expression correlated inversely with the siROS. Further, through experiments with ROS quenchers, we showed that ROS regulated σ^B expression under shear.

Introduction

Hydrodynamic or shear stress, at sublytic levels, significantly affects the macro-level cell responses such as morphology, growth, and productivity (1–5). The macro-level responses result from altered genetic-level responses to the shear stress (6). Although shear is normally perceived to be deleterious, it can also be beneficial (7, 8). Since shear affects cells at a fundamental level, a better understanding of its effects on cells in culture could lead to the design of appropriate bioreactors, or the operation of existing bioreactors in favorable regimes.

Studies on the effects of and cell responses to shear are available (6, 9–11) predominantly on mammalian, plant and insect cells, and molds. However, information on shear responses of bacterial cells, which are widely used in a variety of industries, including those producing high-value products such as pharmaceuticals and other recombinant products, is not as abundant (few studies exist, e.g., 7, 8, 12–14); the information on genetic-level responses of bacterial cells to shear is not available.

Many studies on shear effects have been reported in stirred tank bioreactors, in which it is difficult to decouple

the effects of shear and oxygen transfer. For example, increase in rpm to increase shear rate would also result in an increase in the volumetric oxygen transfer coefficient. Further, it is difficult to quantify the shear levels in a stirred tank, and the average shear rate may not be representative of the actual range of shear levels to which the cells are exposed in the vessel. However, shear studies in the couette flow regime, which is achieved in the space between two concentric cylinders rotating at different angular velocities are available. Although the defined and uniform shear field in couette flow facilitates analysis, it is nontrivial to supply oxygen to aerobic cultures at adequate rates without affecting the flow field for the entire duration of growth. One is therefore constrained to device short-time experiments (e.g., 3), or to introduce the oxygen/air in a manner that might compromise the defined shear field. In their studies on *A. niger* for example, Mitard and Riba (1988) provided oxygen by sparging air into the annular space of their couette flow bioreactor. The effect of such sparging on the flow field is uncertain.

In this work, a couette-flow bioreactor (CFB) has been designed and fabricated to cultivate cells under well-defined laminar shear conditions for the entire duration of growth. The CFB was designed to ensure operation in the couette-flow regime (by using a thin annulus) as well as to ensure oxygen supply without compromising the laminar flow profiles, so that long-term cultivations of

* To whom correspondence should be addressed. Phone: +91 (22) 2576 7208. Fax: +91 (22) 2572 6895/3480. E-mail: gksuresh@che.iitb.ac.in.

[†] Department of Chemical Engineering.

[‡] School of Biosciences and Bioengineering.

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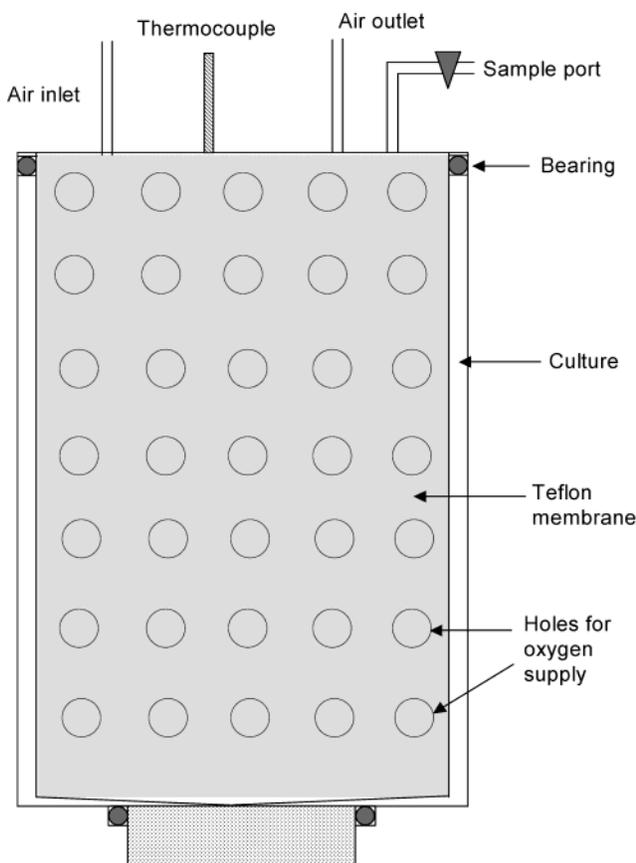


Figure 1. A schematic diagram of the couette flow bioreactor (CFB) with a provision to supply oxygen without disturbing the flow pattern.

an aerobically growing organism can be carried out in a defined shear environment. The problem of oxygen supply with minimum disturbance to the flow field was addressed by having oxygen inside the hollow inner cylinder and transferring it to the annular space containing the culture across an oxygen-permeable membrane.

Using the CFB, the responses of *Bacillus subtilis* cells when grown under well-defined laminar shear conditions were studied. *Bacillus subtilis* was chosen as the model system, because it is one of the preferred hosts in bioindustry and has been well studied. Also, it is nonpathogenic, relatively simple to cultivate, and secretes a variety of proteins and metabolites. In the sections to follow, analyses of the shear rate and mass transfer aspects in the CFB and their comparison with shake flask show that the shake flask can be used as the “control” for shear studies. The effect of shear on the macro-level aspects such as morphology, growth, and enzyme productivity are presented. Also, since reactive oxygen species (ROS) are suspected to be the mediators of many stresses (15), the effect of shear on specific intracellular reactive oxygen species level (siROS) was studied. Further, the involvement of the transcription factor, σ^B , in the shear response of cells was explored. Also, the relationship between (siROS) and expression of σ^B was investigated.

Materials and Methods

The cultivations of *B. subtilis* were carried out in two types of devices in the present work. The first was a standard shake flask of 500 mL capacity, while the other was the CFB designed and fabricated as a part of this work.

Couette Flow Bioreactor (CFB). Figure 1 shows a schematic of the CFB, fabricated at Vaspan Industries,

Mumbai, India. The bioreactor has the inner cylinder stationary and the outer cylinder rotating, a configuration that stabilizes the flow and allows operation at higher Reynolds numbers (16) as compared to the alternative configuration of inner cylinder rotating and the outer stationary. The reactor has been designed to be able to accommodate a culture volume of 250 cm³ in the annular space. Care has been taken to see that the thickness of the annular region is small compared to the outer radius, the ratio of radii of the inner to the outer cylinder being about 0.93. Similarly, the height occupied by the culture volume in the annulus is sufficiently large as compared to the thickness of the annular region (the ratio being 76.05) so that end effects are negligible. A uniform shear field can therefore be assumed to exist in the annular space when the device is operated.

The inner cylinder is hollow, and a regular square array of 10 mm diameter holes is made in the curved surface to provide the mass transfer area for oxygen. A poly(tetrafluoroethylene) (PTFE) membrane is stuck over this surface, with care taken to smooth out any wrinkles. A port is provided to admit air into the hollow space inside the inner cylinder, and oxygen from this space diffuses across the membrane and is transferred into the culture by a process of convective diffusion.

Ports are provided for sample withdrawal and for a thermocouple, with care taken care not to interfere with the flow pattern. The entire assembly is housed in a Plexiglas enclosure so that the desired temperature can be maintained by controlling the temperature of air in the enclosure. The whole reactor, with the membrane in place, is autoclaved prior to cultivation, and the culture is transferred under the sterile conditions provided by a laminar hood.

Cell Strains, Media, and Cell Treatment. *Bacillus subtilis* 168 (*trpC2*) (BGSC, Ohio State University, Columbus, OH) cultures were stored at -20 °C in LB liquid media (Hi-media) containing 15% glycerol. The *sigB* null-mutant of *B. subtilis* (*sigB::ΔHindIII-EcoRV::cat trpC2*, BSA272, (17)) was kindly provided by Prof. W. Haldenwang, Department of Microbiology, University of Texas, San Antonio, TX. The *sigB-lacZ* strain of *Bacillus subtilis* (*amyE::sigB-lacZ trpC2*, PB286, (18)) was a kind gift from Prof. C. W. Price, Food Science Department, University of California, Davis, CA. PB286 carries a transcriptional fusion of σ^B to the *E. coli lacZ* gene. This fusion allows β -galactosidase activity to be monitored as a measure of σ^B expression.

The cells were grown in Luria-Bertani (LB) medium (Hi-media). The preinoculum was prepared by growing cells in liquid media at 37 °C to exponential growth phase (12 h). They were diluted 1:10 (v/v) into the fresh media and inoculated either to the CFB or to the shake flasks. The initial cell concentration was maintained at 0.5 ± 0.03 g L⁻¹ by adjusting the preinoculum concentration. Special care was taken for transferring the preinoculum to the autoclaved CFB, inside a laminar flow hood. The cells were grown in different shear environments (in the CFB at various rpm or, in the shake flasks, at 190 rpm) at 37 °C for 30 h. All experiments were repeated at least twice for reproducibility. The maximum variation in corresponding data points was 9%.

The antioxidant *N*-acetyl cysteine (NAC), as well as diphenylene iodoniumchloride (DPI), an inhibitor of superoxide generation, were used as treatment agents in certain studies. The concentrations of NAC and DPI used were 5 mM and 0.6 μ M, respectively. The concentrations of NAC and DPI used were determined through toxicity studies; cells were exposed to different initial

Table 1. Estimated Volumetric Oxygen Transfer Coefficients (k_{LA}) in CFB without and with PTFE Membrane on Inner Cylinder, Calculated Value of k_{LA} in Shake Flask, Shear Rates at Various rpm, and Specific Growth Rates for Different Cultivations

rpm	estimated k_{LA} (without membrane) (h^{-1})	measured/estimated k_{LA} (with membrane in CFB) (h^{-1})	shear rate (s^{-1})	specific growth rate ^a (h^{-1})
(CFB) 300	256	4.54	445	0.26
(CFB) 500	407	4.56	741	0.33
(CFB) 750	554	4.58	1111	0.40
(CFB) 1000	738	4.59	1482	0.76
(shake flask) 190	na	2.89	0.028	0.22

^a Each specific growth rate value was obtained from the average values of cell concentrations from two/three different experiments at each shear rate.

184 concentrations of the two treatment agents, and the
185 concentration that least affected the viability was chosen
186 for treatment. NAC was added to the medium at the start
187 of the cultivation, and cells were exposed to DPI (pre-
188 pared fresh) for 5 min, separated from the treatment
189 medium, resuspended in cell-free medium of similar
190 nature, and grown.

191 **Analyses.** Samples were taken from the shake flask
192 and the CFB at regular intervals. Each sample of
193 approximately 5 mL was collected, and cells were sepa-
194 rated from the supernatant (centrifugation at 12,000 \times
195 g for 15 min). They were stored in a deep freezer (-20
196 $^{\circ}\text{C}$) until further analysis was done. For measuring
197 intracellular ROS levels, the sample was processed
198 immediately. Cell concentration was measured using a
199 spectrophotometer (Shimadzu, Japan) through cell scat-
200 ter measurements at 600 nm and using a calibration
201 graph. Standard methods were followed for measuring
202 extracellular enzyme concentrations immediately after
203 the completion of the cultivation: the azocasein degrada-
204 tion method for protease (19), the hydrogen peroxide
205 decomposition rate method for catalase (20), and the
206 starch hydrolysis method for amylase (21). The standard
207 curve was prepared using enzymes from Sigma. A
208 standard assay procedure for β -galactosidase (22) was
209 followed, and specific activity was expressed in Miller
210 Units (MU), which was calculated as

$$\text{MU} = \frac{A_{420} \times 1000}{A_{600} \times \text{reaction time in min}}$$

211 **Spin Labeling and Electron Spin Resonance**
212 **(ESR) Spectroscopy.** One milliliter of the fresh sample
213 was centrifuged (12,000 $\times g$, 10 min, -4°C) and washed
214 thrice with saline to remove traces of media with extra-
215 cellular secretions. The cells were suspended with an
216 equal amount of 100 mM ice-cold Tris buffer (pH 8.0,
217 saturated with N_2). Lysozyme 2.5 mg mL^{-1} , sucrose 100
218 mM, EDTA 132 μM , and MgCl_2 2.5 mM were added, and
219 the mixture was incubated at 37 $^{\circ}\text{C}$ for 15 min. The
220 solution was centrifuged at 12,000 $\times g$, 4 $^{\circ}\text{C}$ for 10 min.
221 The resulting protoplasts in the pellets were washed once
222 with the above mixture without lysozyme. To the proto-
223oplasts was added 80 mM dimethyl-1-pyrroline-*N*-oxide
224 (DMPO, Sigma) (spin trap). Subsequently, the volume
225 was made up to 1 mL using distilled water and mixed
226 thoroughly to break open the protoplasts through osmotic
227 shock. The derivative ESR spectrum was obtained im-
228 mediately after, using a Varian E-112 EPR spectrometer.
229 The final two steps were done under red light. The area
230 under the absorption curve was obtained by double
231 integration of the derivative ESR spectrum using a
232 computer program, and a standard curve was used to
233 obtain the actual ROS concentrations; the free radical
234 concentration is proportional to the area under the
235 absorption curve (23). The specific intracellular ROS level

236 was obtained by normalizing the intracellular ROS
237 concentrations with the corresponding viable cell number
238 concentrations. The viable cell number concentration was
239 determined by the plating method, and the plates were
240 inoculated immediately after sample collection. The ROS
241 type was identified by comparison with standard spectra
242 in the literature (24).

243 **Sample Preparation for Scanning Electron Mi-**
244 **croscopy (SEM).** The SEM technique was used to
245 examine the cell shape and size at high resolution. Fresh
246 samples of cells taken from the CFB or shake flasks were
247 used for analysis using a Quanta 200, FEI-Philips
248 environmental scanning electron microscope (ESEM). For
249 analysis using ESEM, a carbon-coated copper grid was
250 dipped in the sample for 1 min and then dipped in 0.5%
251 phospho-tungstic acid for 1 min for negative staining of
252 the cells. They were dried with the help of Whatman No.
253 1 filter paper. The grids were stuck to a stub and placed
254 on the specimen holder under vacuum. The cells were
255 viewed under low vacuum mode at room temperature
256 under different magnifications and acceleration voltages.

257 Results and Discussion

258 **Oxygen Transfer Coefficient in CFB.** Although the
259 CFB has provision for supplying oxygen without actual
260 sparging and thereby not interfering with the flow field,
261 it is important to make sure that the transfer rates
262 provided for are adequate. An estimate of the mass
263 transfer rate was obtained by measuring the volumetric
264 mass transfer coefficient k_{LA} by a physical method as
265 follows: the inner cylinder, with air passing in and out
266 to maintain a constant oxygen partial pressure, was held
267 in a vessel containing water depleted of oxygen by prior
268 sparging of nitrogen. Oxygen tension in the water was
269 measured as a function of time until saturation was
270 reached. Oxygen transfer to the water in the vessel
271 through other routes such as that across the air–water
272 interface on the top of the vessel was avoided by using
273 aluminum foil to cover the surface. A semilog plot of
274 fractional saturation vs time yielded a value of 0.17 h^{-1}
275 for k_{LA} , under quiescent conditions.

276 The values of the volumetric oxygen transfer coef-
277 ficient, k_{LA} , in the CFB under convective conditions were
278 estimated as follows. From a plot of j_D' vs Re for mass
279 transfer in a rotating cylinder system (25) the k_{LA} values
280 were estimated using the appropriate values for angular
281 velocities, Schmidt numbers, and transfer areas per unit
282 volume. The k_{LA} values, given in Table 1, show that, as
283 expected, under convective conditions they were 3 orders
284 of magnitude higher than those under the quiescent
285 conditions. However, in the system employed here, the
286 gas and the liquid phase are separated by the PTFE
287 membrane, and it is important to account for the trans-
288 port resistance due to the membrane. This can be
289 accomplished by writing a quasi steady-state equation
290 for the oxygen mass transfer flux N_{O_2} , considering the

D

291 transport processes in series:

$$N_{O_2} = \left(\frac{D}{\delta} \right) (H_1 p_{O_2} - c_\delta) = k_L (H_2 c_\delta - c_b) \quad (1)$$

292 where p_{O_2} is the partial pressure of oxygen inside the
 293 inner cylinder, D is the diffusivity of oxygen through the
 294 membrane, δ is the membrane thickness, c_δ is the
 295 concentration of oxygen in the membrane phase at the
 296 membrane–liquid interface, and c_b is the concentration
 297 of oxygen in the liquid bulk. H_1 is the Henry’s law
 298 coefficient for oxygen for the gas–membrane system, and
 299 H_2 is the partition coefficient for oxygen between the
 300 membrane and the liquid. A solution-diffusion process
 301 for the transport across the membrane has been assumed,
 302 and transport resistance on the gas side has been
 303 neglected. Knowing one of H_1 and H_2 and the Henry’s
 304 law coefficient H for oxygen for the air–water system,
 305 the other can be calculated, as the equilibrium between
 306 air and water should not depend on whether a membrane
 307 separates the phases:

$$H = H_1 H_2 \quad (2)$$

308 Calculating c_δ from eq 1 and substituting in one of the
 309 right-hand sides of the same equation enable the mass
 310 transfer flux to be written in terms of an overall liquid
 311 side driving force:

$$N_{O_2} = k_{L,\text{eff}} (H p_{O_2} - c_b) \quad (3)$$

312 where

$$k_{L,\text{eff}} = k_L \frac{\frac{D}{\delta}}{\frac{D}{\delta} + k_L H_2} \quad (4)$$

313 $k_{L,\text{eff}}$ is the effective overall mass transfer coefficient
 314 that accounts for the resistances due to the membrane
 315 as well as the liquid film. Clearly, the importance of the
 316 membrane resistance depends on the thickness of the
 317 membrane for a given material. For a given membrane,
 318 the importance of its resistance, vis-à-vis the liquid side
 319 resistance, depends on the value of the latter; the
 320 membrane resistance increases in importance as the
 321 liquid side resistance decreases. For the PTFE membrane
 322 used, we have the following data at 25 °C (the diffusivity
 323 and solubility of oxygen in PTFE have been taken from
 324 <http://www.sablesys.com/oxelect.html> accessed on Sep-
 325 tember 10, 2002): thickness (δ) = 100 μm ; $D = 2.54 \times$
 326 $10^{-11} \text{ m}^2 \text{ s}^{-1}$; $H_1 = 0.106 \times 10^{-7} \text{ mol (m}^3 \text{ kPa)}^{-1}$; and H
 327 $= 1.19 \times 10^{-8} \text{ mol (m}^3 \text{ kPa)}^{-1}$ (26). Substitution of these
 328 values gives

$$k_{L,\text{eff}} = k_L \frac{2.54 \times 10^{-7}}{2.54 \times 10^{-7} + 1.12 k_L} \quad (5)$$

329 Values of the effective overall volumetric mass transfer
 330 coefficient $k_{L,\text{eff}} a$ calculated from the above equation,
 331 using the estimated k_L values for different rpm’s in the
 332 CFB are shown in Table 1, and it is clear that the
 333 membrane resistance dominates. As a consequence, ef-
 334 fective transfer rates are nearly constant across all of the
 335 rotation speeds used. Significantly, in the range of rpm
 336 employed, the $k_{L,\text{eff}} a$ value is between 4.54 and 4.59 h^{-1} ,
 337 which is of the same order of magnitude as that measured
 338 in the shake flask at 190 rpm (2.89 h^{-1}). Thus CFB is an

339 ideal device to study the effect of shear alone, which is
 340 not possible in stirred tanks; in stirred tanks the com-
 341 bined effects of shear and oxygen transfer cannot be
 342 easily decoupled when air is used as the oxygen source.

Shake Flask Cultivations Used as Control Cultivations for Shear Studies. Two types of devices have
 343 been used in the present cultivations, the couette flow
 344 bioreactor (at several speeds of rotation to vary the shear
 345 rate) and the shake flask. For a meaningful comparison
 346 of the results in the two devices, we need methods of
 347 calculating the shear rates in the two devices. The shear
 348 rates in the CFB can be exactly calculated from the
 349 velocity field (27, 28):
 350
 351

$$v_\theta(r) = R\Omega \frac{r/(kR) - kR/r}{(1/k) - k} \quad (6)$$

352 where k is the ratio of the inner radius to the outer
 353 radius R , and Ω is the angular velocity of the outer
 354 cylinder. Differentiating, we get the shear rate at any
 355 radial position r as

$$\dot{\gamma}(r) = 2\Omega \frac{\left(\frac{kR}{r} \right)^2}{k^2 - 1} \quad (7)$$

356 For thin annuli ($k \rightarrow 1$), these equations predict a
 357 nearly linear profile of v_θ , and hence, a nearly constant
 358 shear rate at every r .

359 For the considerably more complex and turbulent
 360 hydrodynamics prevailing in the shake flask, the shear
 361 rate can only be estimated. The energy dissipation rate
 362 in shake flasks is reasonably uniform (especially in
 363 comparison with mechanically stirred bioreactors), so
 364 that an average shear rate can be used for purposes of
 365 comparison (28). Extending the same argument further,
 366 such an average shear rate can be calculated for the
 367 shake flask in the following manner.

368 The shear stress on a particle at any location in a
 369 turbulent flow field is determined primarily by the
 370 velocity of turbulent fluctuations. The theory of isotropic
 371 turbulence visualizes eddies of a spectrum of sizes in a
 372 turbulent flow field, the energy content of the eddies
 373 decreasing with their size. In such a scenario, the stress
 374 that a particle feels in the flow field depends on the
 375 particle size relative to an appropriate length scale of
 376 turbulent eddies. Kolmogorov’s microscale of turbulence
 377 η_1 is normally used as a measure of the latter and is
 378 related to the rate of energy dissipation ϵ as follows:

$$\eta_1 = \left(\frac{V^3}{\epsilon} \right)^{1/4} \quad (8)$$

379 For the shake flask, since the rates of energy dissipation
 380 do not vary greatly across the cross section, we can
 381 associate a Kolmogorov length scale with the *average* rate
 382 of energy dissipation, calculated as the power input per
 383 unit mass. The power input for shake flasks can be
 384 calculated from (28):

$$\frac{P}{\rho n^3 d_s^5} = \frac{1.94}{Re^{0.2}} \frac{V_L^{1/3}}{d_s} \quad (9)$$

385 where the Reynolds number Re is defined as

$$Re = \frac{nd_s^2}{\nu} \quad (10)$$

386 For the conditions relevant to our experiments, these
 387 equations give a value of $4.32 \times 10^{-5} \text{ m}$ for η_1 . Electron
 388 micrographs (discussed later) show the *Bacilli* grown in
 389 shake flasks to be about $1 \mu\text{m}$ in diameter and $3\text{--}4 \mu\text{m}$
 390 in length. Since the particle dimension is almost an order
 391 of magnitude smaller than η_1 , we may expect the effect
 392 of shear to be negligible in the shake flask. For such small
 393 ratios of particle diameter to the Kolmogorov scale, the
 394 determining eddies are in the dissipation range, for which
 395 the shear stress τ_t is given by the equation (28)

$$\frac{\tau_t}{\rho\sqrt{v\epsilon}} = 0.0676\left(\frac{d_p}{\eta_1}\right)^2 \quad (11)$$

396 Choosing a spherical diameter d_p of $2 \mu\text{m}$ for the
 397 *Bacillus* for an estimate, we get the shear stress to be
 398 $2.8 \times 10^{-5} \text{ N m}^{-2}$. For water-like viscosity, this would
 399 correspond to a shear rate of about 0.028 s^{-1} . This value
 400 is very small, particularly when compared with the
 401 values of shear rate that can be calculated for the CFB,
 402 confirming the expectation that the *Bacilli* experience
 403 negligible shear in the shake flask. These calculations,
 404 together with those in the previous section, which show
 405 the oxygen transfer rates in the two devices to be similar
 406 in order of magnitude, therefore suggest that experiments
 407 in the shake flask can be regarded as control experiments
 408 in delineating the effects of shear on fermentation
 409 behavior.

410 **Cell Size.** The cell size under shear was studied using
 411 scanning electron microscopy. The average cell size
 412 obtained at 0.028 s^{-1} was about $3.05 \mu\text{m}$ and reduced to
 413 about half that size when grown at 1482 s^{-1} in the CFB
 414 (Figure 2). This was in contrast to our initial expectation
 415 of longer cells at higher shear rates, as observed with
 416 bacterial cells exposed to short duration of laminar shear
 417 (7). In any case, those authors suspected that the cell
 418 elongation was due to the added viscosity modifier.
 419 Probably, the size reduction observed in this study is due
 420 to cell adaptation mechanisms when exposed to a higher
 421 shear rate. For example, it may be an advantage in terms
 422 of stress handling for the cells to present a smaller
 423 surface to the flow around them when the levels of shear
 424 stress are high. In addition, asymmetrical cell division
 425 was noticed at shear rate 1482 s^{-1} ; some dividing cells
 426 appeared pinched off at one end. Similar asymmetrical
 427 cell divisions were observed by Edwards et al. (1989) in
 428 *E. coli* cells at high shear levels (10.9 and 14.5 N m^{-2}).

429 **Growth and Enzyme Production.** The time profiles
 430 of cell concentration obtained at different shear rates are
 431 presented in Figure 3; the average shear rates and the
 432 corresponding rpm are presented in Table 1. Interest-
 433 ingly, the maximum cell concentration obtained increased
 434 with the shear rate. The maximum cell concentration
 435 obtained at the highest average shear rate studied, 1482
 436 s^{-1} (1000 rpm), was 7.8 g L^{-1} , which was 195% of that
 437 obtained at 445 s^{-1} (300 rpm) and 289% of that obtained
 438 at 0.028 s^{-1} (shake flask control at 190 rpm). The
 439 exponential specific growth rates, presented in Table 1,
 440 show that it also increased significantly with shear rate;
 441 the exponential specific growth rate at 1482 s^{-1} was 292%
 442 of that obtained at 445 s^{-1} and 345% of that obtained at
 443 0.028 s^{-1} . Also, the cells grown at higher shear rates
 444 showed a shorter lag period as compared to the cells
 445 grown at 0.028 s^{-1} . The above observations that the
 446 exponential specific growth rate and maximum cell
 447 concentration increase, whereas the lag times decrease
 448 with increase in shear rate, can be utilized to reduce the
 449 cultivation time significantly in industrial bioreactors,

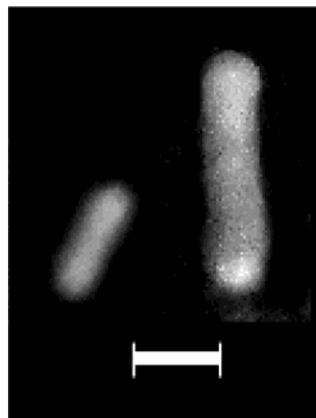


Figure 2. Scanning electron micrographs of cells grown in a shake flask at a shear rate of 0.078 s^{-1} (right) and in CFB at a shear rate of 1482 s^{-1} (left). ESEM, $8000\times$, 20 kV , 0.98 Torr ; bar = $1 \mu\text{m}$.

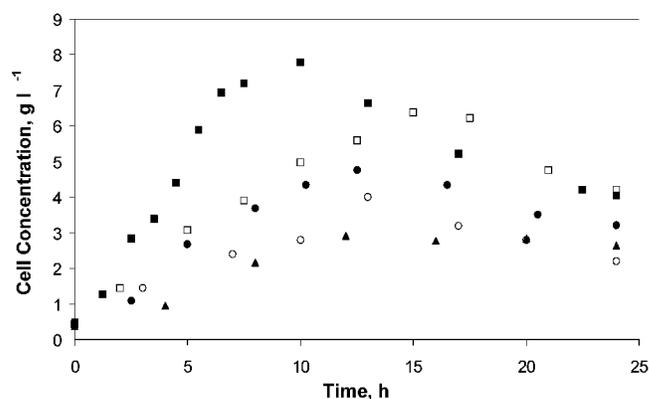


Figure 3. Growth profiles under various shear rates: (■) 1482 , (□) 1111 , (●) 741 , (○) 445 , (▲) 0.028 s^{-1} .

which would improve the process economics; however, the
 economical achievement of higher shear rates could be a
 challenge.

The maximum specific levels of extracellular enzymes,
 i.e., catalase, protease, and amylase, obtained at various
 shear rates are presented in Figure 4. From the figure,
 it can be seen that the specific levels of catalase increased
 with shear rate: the maximum specific level at 1482 s^{-1}
 was $3035 \text{ U (g-cell)}^{-1}$, which was 199% of that obtained at
 445 s^{-1} and 765% of that obtained at 0.028 s^{-1} . The
 extracellular protease level also increased with shear
 rate: the maximum specific level at 1482 s^{-1} , 0.73 U (g-
 $\text{cell)} \text{ L}^{-1}$, was 252% of that obtained at 445 s^{-1} and 187%
 of that obtained at 0.028 s^{-1} . The reason for the initial
 decrease in specific protease level with shear rate is
 unclear. However, the extracellular specific levels of
 amylase did not vary significantly with shear rate in the
 range studied.

Correlation between Shear and siROS. Reactive
 oxygen species are known to mediate cellular processes
 under many stress conditions (15). Also, ROS are sus-
 pected to be involved in mediating shear stress responses
 of animal cells (29, 30). Therefore, we suspected that ROS
 are involved in shear stress responses of bacteria also,
 and the specific intracellular ROS levels were determined
 when the wild-type cells were grown at various shear
 rates. The ROS type was found out to be superoxide and
 superoxide-derived radicals. The 10-h siROS presented
 in Figure 5 clearly show that the siROS increased with
 shear rate; it was $3.7 \text{ mmol (g-cell)} \text{ L}^{-1}$ at 1482 s^{-1} , which
 was 9.3-fold of that at 0.028 s^{-1} and 4.6-fold of that at

F

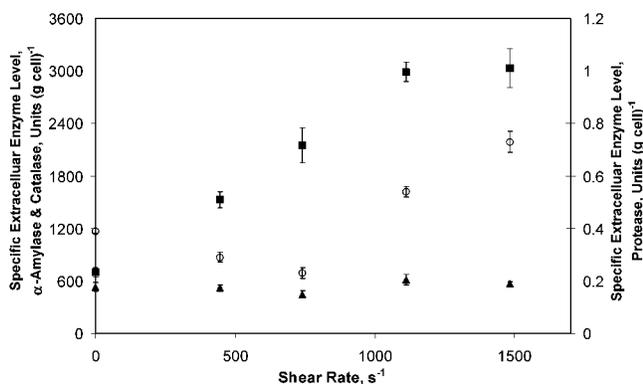


Figure 4. Maximum specific levels of (■) extracellular catalase, (○) protease, and (▲) α -amylase at different shear rates. The specific enzyme levels were determined by normalizing the enzyme activity with the cell concentration, at the corresponding time.

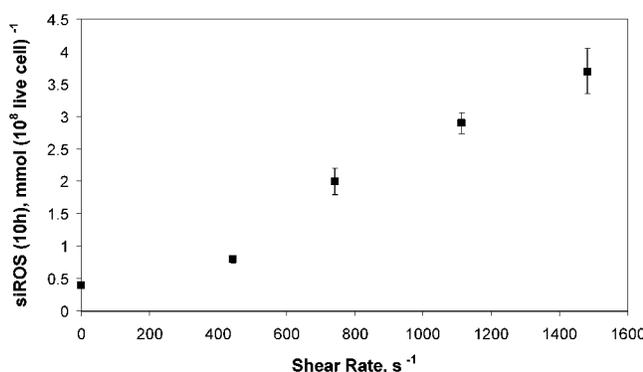


Figure 5. Specific intracellular reactive oxygen species levels (siROS) obtained at 10 h (maximum siROS) when cultivated at various shear rates.

445 s⁻¹. The siROS at other times also showed the same trend (data not shown). Therefore, it is reasonable to suggest that ROS are involved in mediating shear stress responses in *B. subtilis*.

Genetic Basis for Shear Response; Involvement of σ^B . To understand the different responses of the cells under shear stress, the general stress transcription factor, σ^B was chosen for the study. σ^B -controlled genes are known to play important roles in protection against the deleterious effects of various stresses such as oxidative, heat shock, acid, ethanol, and salt stresses and nutrient depletion (31), although the relationship between shear stress and σ^B is unknown thus far. Hence, we were interested in determining if σ^B also played a role in responses to shear stress. In addition, *katE*, which is one of the genes responsible for catalase production in the stationary phase, is controlled by σ^B at the transcription level (32). Therefore, we expected a reduction in the stationary-phase catalase level when σ^B was inactive.

To investigate the above possibilities, a strain that lacks σ^B (*sigB* null-mutant, *sigB::cat trpC2*) was compared with the wild-type strain, which produces σ^B (*trpC2*), at the highest shear rate, 1482 s⁻¹. From the results presented in Table 2, it can be seen that the specific extracellular catalase level obtained with *sigB* null-mutant, *sigB::cat*, was 60% lower than that obtained with the wild-type strain. In addition, the maximum cell concentration obtained with *sigB::cat* was 25% lower than that obtained with the wild-type, indicating that the *sigB* null-mutant was more shear-sensitive compared to the wild-type strain. Under low shear conditions (0.028 s⁻¹), the catalase productivity and growth of the *sigB* null-

Table 2. Comparison of Maximum Specific Enzyme Levels, Maximum Cell Concentrations, and Specific Growth Rates Obtained in Cultivations at 1482 s⁻¹ of *sigB*-null Mutant and Wild-Type Cells

parameter	wild-type	<i>sigB</i> -null mutant
maximum specific catalase level, Units (g cell) ⁻¹	3035 ± 202	1229 ± 59
maximum specific amylase level, Units (g cell) ⁻¹	570 ± 22	368 ± 24
maximum specific protease level, Units (g cell) ⁻¹	0.73 ± 0.04	1.25 ± 0.08
maximum cell concentration, g L ⁻¹	7.8 ± 0.4	5.9 ± 0.5
specific growth rate, h ⁻¹	0.76	0.74
max siROS, mmol (10 ⁸ live cells) ⁻¹	3.7 ± 0.4	11.3 ± 0.6

mutant was comparable to those of the wild-type cell. Therefore, it is clear that σ^B is involved in responses to shear stress and possibly protects the cells under high shear.

Interestingly, at 1482 s⁻¹ the specific extracellular amylase level with the *sigB* null-mutant was 35% lower than that with the wild-type; the growth rates of both were comparable, and the specific extracellular protease level with the *sigB* null-mutant was 71% higher than with the wild-type; the reasons for these observations are unclear.

To further investigate the relationship between shear and σ^B , we measured the transcriptional levels of σ^B in the strain *sigB::lacZ trpC2*. The data given in Table 3 shows that the expression of σ^B decreased with increasing shear rates. For example, the maximum transcription level of σ^B at 445 s⁻¹ was 42% lower, and at 1482 s⁻¹ it was almost 40-fold (4000%) lower, compared to 275 MU at 0.028 s⁻¹. The maximum expression of σ^B was obtained in the late-exponential/stationary phase (data not shown); the zero time values were not considered because they represent the inoculum conditions. The finding that σ^B protects the cells even when maintained low at high shear stress can be explained on the basis of the intracellular pool of σ^B that was carried from the inoculum; the pool may remain active and provide the protection. The pool diminishes afterward, because new σ^B molecules are not produced under high shear. In the case of the mutant, the pool is nonexistent even in the inoculum, and therefore these cells are more sensitive to the shear stress. However, the reasons for this decreased expression of σ^B in a highly stressed cell, where it is shown to protect the cell, is still unclear. One of the reasons could be the unusually high level of intracellular ROS (superoxide and superoxide-derived radicals) under shear stress.

Relationship between ROS and σ^B under Shear Stress. To investigate the possible relationship between the stress mediator, ROS, and σ^B expression under shear stress, we measured the siROS and the σ^B expression at very low (0.028 s⁻¹), moderate (445 s⁻¹), and high (1482 s⁻¹) shear levels. The composite data in Figure 6 shows that σ^B transcription level decreased with increase in siROS. It decreased steeply from a value of around 260 MU at 0.4 mmol (g-cell) L⁻¹ to about 15 MU at 1 mmol (g-cell) L⁻¹. It reached a basal value of about 6 MU at 2 mmol (g-cell) L⁻¹ siROS.

To further support the relationship between ROS and σ^B expression under shear, we studied the transcription level of σ^B when siROS was reduced at the highest shear rate employed, 1482 s⁻¹. The antioxidant, *N*-acetyl cysteine (NAC), was used to reduce siROS. In the cultivation with NAC, the maximum siROS was 0.9 mmol (g cell) L⁻¹, which was 4-fold lower compared to the cultivation without NAC. The corresponding σ^B level with NAC was

Table 3. Maximum Specific Intracellular σ^B Transcription Levels As Indicated by β -Galactosidase, Obtained at 13 h, during Cultivation at Various Shear Levels^a

shear rate, s ⁻¹	maximum specific intracellular transcription level of <i>sigB</i> (MU)
0.028	275 ± 16
445	127 ± 11
1482	7 ± 0.5
1482 + DPI	162 ± 12

^a The NADH-oxidase (superoxide generator) inhibitor, DPI, was added at 9 h after the start of cultivation in CFB.

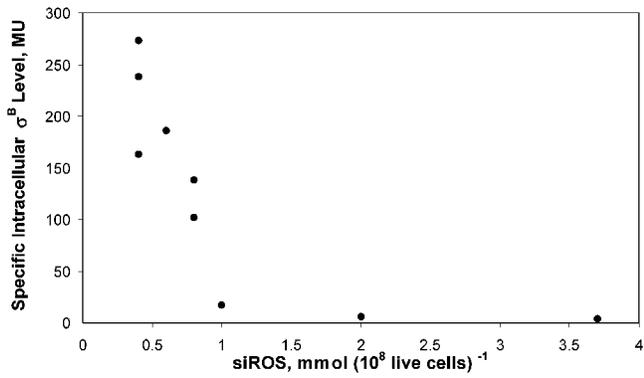


Figure 6. Specific intracellular σ^B transcription levels as indicated by β -galactosidase, in Miller Units (MU), at various specific intracellular ROS level (siROS) obtained in different cultivations.

567 13 MU, which was about 2-fold higher compared to the
 568 cultivation without NAC. Further, when diphenylene
 569 iodoniumchloride (DPI), an inhibitor of ROS generating
 570 plasma membrane protein NADH-oxidase (NOX) was
 571 used to reduce siROS to a basal level (0.4 mmol (g cell)
 572 L⁻¹), the σ^B expression increased by about 10-fold (Figure
 573 3). These results clearly indicate that σ^B expression under
 574 shear is regulated by intracellular levels of superoxide
 575 radical.

576 In addition, the siROS obtained in the σ^B null mutants
 577 was compared with those obtained in wild-type cells
 578 under high shear (1482 s⁻¹) conditions. The results
 579 presented in Table 2 show that the maximum siROS in
 580 the absence of σ^B was almost 3-fold higher compared to
 581 those obtained when σ^B was present. This indicates that
 582 σ^B can reduce the iROS level under high shear conditions.
 583 One of the ways in which σ^B reduces iROS level could be
 584 by regulating the antioxidant enzymes that quench ROS,
 585 such as catalase; we had seen earlier that catalase
 586 expression is controlled by σ^B .

587 A possible mechanism of ROS and σ^B involvement in
 588 determining the cell responses to shear stress is sche-
 589 matically shown in Figure 7.

Conclusions

591 Under defined shear stress conditions, the growth,
 592 morphology, and enzyme productivity of *Bacillus subtilis*
 593 and, more importantly, its genetic responses were stud-
 594 ied. A couette flow bioreactor with a capability to provide
 595 oxygen over the entire cultivation period without affect-
 596 ing the laminar flow profile was used for the study; shake
 597 flask cultivations were established as the control cultiva-
 598 tions. Shear stress significantly affected growth charac-
 599 teristics, morphology, and enzyme productivity of *B.*
 600 *subtilis*. More interestingly, the shear level influenced
 601 gene expression. The stress transcription factor σ^B was
 602 involved in cell responses to shear. Also, the specific

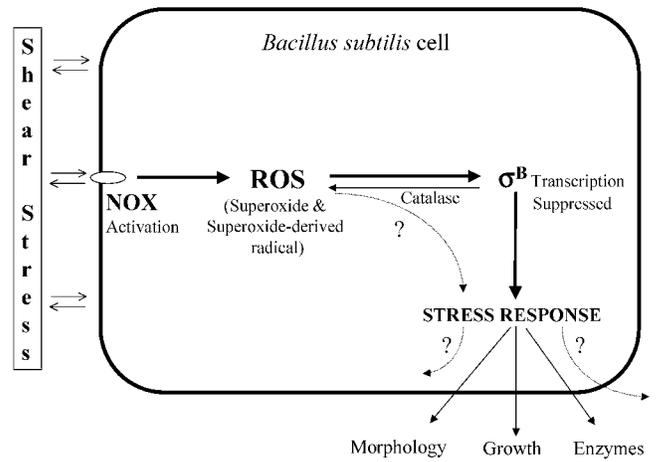


Figure 7. A schematic representation of some of the intracellular effects of shear stress in *Bacillus subtilis*. NOX: NADH oxidase. ROS: reactive oxygen species. σ^B : transcription factor.

intracellular reactive oxygen species level (siROS) in- 603
 creased with shear; the siROS and σ^B expression under 604
 shear are interrelated. 605

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 mentation Center. 613

Notation

c_b	oxygen concentration in the liquid bulk (mol m ⁻³)	615
c_δ	oxygen concentration at the membrane-liquid bound- 616 ary (mol m ⁻³)	617
d_p	particle (cell) diameter (m)	618
d_s	diameter of the rotating liquid mass (m)	619
D	diffusivity of oxygen through the membrane (m ² s ⁻¹)	620
H	Henry's law coefficient for oxygen, gas/liquid system 621 (mol (m ³ N m ⁻²) ⁻¹)	622
H_1	Henry's law coefficient for oxygen, gas/membrane 623 system (mol (m ³ N m ⁻²) ⁻¹)	624
H_2	Henry's law coefficient for oxygen, membrane/liquid 625 system (mol (m ³ Nm ⁻²) ⁻¹)	626
k_L	liquid side mass transfer coefficient (m s ⁻¹)	627
$k_{L,eff}$	effective liquid side mass transfer coefficient (eq 4) 628 (m s ⁻¹)	629
n	rotational speed (s ⁻¹)	630
N_{O_2}	oxygen mass transfer flux (mol (m ² s ⁻¹) ⁻¹)	631
p_{O_2}	partial pressure of oxygen in the inner cylinder (N 632 m ⁻²)	633
P	power input (shake flask) (W)	634
r	any radial position in the annulus of CFB (m)	635
R	radius of the outer cylinder (m)	636
Re	Reynolds number (as defined in eq 10)	637
$v_\theta(r)$	tangential velocity of the fluid at the location r (m 638 s ⁻¹)	639
V_L	liquid volume in the shake flask (m ³)	640
Greek letters		641
$\dot{\gamma}$	shear rate at any location in the annulus (s ⁻¹)	642
δ	membrane thickness (m)	643
ϵ	rate of energy dissipation (W kg ⁻¹)	644
η_l	Kolmogorov lengthscales (m)	645

646	κ	ratio of radii of the inner cylinder to the outer cylinder
647	ν	kinematic viscosity ($\text{m}^2 \text{s}^{-1}$)
648	Ω	angular velocity of the outer cylinder (s^{-1})
649	ρ	liquid density (kg m^{-3})
650	τ_t	shear stress (N m^{-2})

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