

Supplementary Information

for

How *Escherichia coli* lands and forms cell clusters on a surface: a new role of surface topography

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Table S1. List of bacterial strains and plasmids used in this study.

| Strain | Genotype | Characteristics | Source/References |
|-------------------------------------|--|---|-------------------|
| <i>E. coli</i> strains | | | |
| RP437 | Wild type [<i>thr-1 leuB6</i> (Am) <i>hisG4</i> (Oc) <i>fhuA31</i> <i>metF159</i> (Am) <i>thiE1 rfbC1 eda-50</i> <i>rpsL136</i> (strR) <i>araC14 mtl-1</i> <i>xylA5 λ⁻ tsx-78 lacY1 F-</i>] | Wild type strain for biofilm study | 1 |
| RP3087 | RP437 (<i>motB</i>)580 | Motility mutant (point mutation in the <i>motB</i> gene) | 2 |
| KX1485 | RP437 Δ <i>luxS</i> ::Cm ^r | Quorum sensing mutant, unable to synthesize AI-2 | 3,4 |
| RHG01 | RP437 Δ <i>fliC</i> ::Kan ^r | Unable to synthesize the subunits of flagella | This study |
| RHG02 | RP437 Δ <i>fimA</i> ::Kan ^r | Unable to synthesize the subunits of fimbriae | This study |
| <i>P. aeruginosa</i> strains | | | |
| PA14 | Wild type [Human clinical isolate; Rif ^r] | Wild type strain for biofilm study | 5,6 |
| PA1092 | PA14 (<i>fliC</i>) | Flagellin type B transposon insertion mutant | 7 |
| PA4526 | PA14 (<i>pilB</i>) | Type 4 fimbrial biogenesis protein PilB transposon insertion mutant | 7 |
| Plasmids | | | |
| pRSH103 | <i>rfp</i> , Tet ^r | To label the cells with constitutive red fluorescence for imaging | 8 |
| pRHG03 | <i>motB</i> ⁺ , Tet ^r | pRSH103 with <i>motB</i> gene controlled by <i>lac</i> promoter | This study |
| pRHG04 | <i>fliC</i> ⁺ , Tet ^r | pRSH103 with <i>fliC</i> gene controlled by <i>lac</i> promoter | This study |
| pRHG05 | <i>fimA</i> ⁺ , Tet ^r | pRSH103 with <i>fimA</i> gene controlled by <i>lac</i> promoter | This study |

Table S2. Primers used for constructing the *E. coli* mutants.

| Target gene | Forward primer (5'-3') | Reverse primer (5'-3') |
|--------------------|-------------------------------|-------------------------------|
| <i>fimA</i> | ACCGTTCACCCGTTACGTTT | TGCTCTTGTTTTTGCCCTGC |
| <i>fliC</i> | CGCCACTCATCGTAGGAGAA | CCGAGACCTCCATACCGTTG |

Table S3. Primers used for complementing the *E. coli* mutants.

| Target gene | Forward primer (5'-3') | Reverse primer (5'-3') |
|--------------------|---------------------------------------|--|
| <i>motB</i> | ATTTGCGGCCGCTGTGCGTGC GGTGAAAAATC | GGACTAGTCATGTCAGCCAACAG TTCGTC |
| <i>fliC</i> | ATTTGCGGCCGCTAGCGGGAA TAAGGGGCAGAG | ATTTGCGGCCGCGAGTCTCAGTTAA TCAGGTTACAACG |
| <i>fimA</i> | ATTTGCGGCCGCGCCAGTAAT GCTGCTCGTTTT | GGACTAGTTTATCGCACAAAGGGT GGGC |

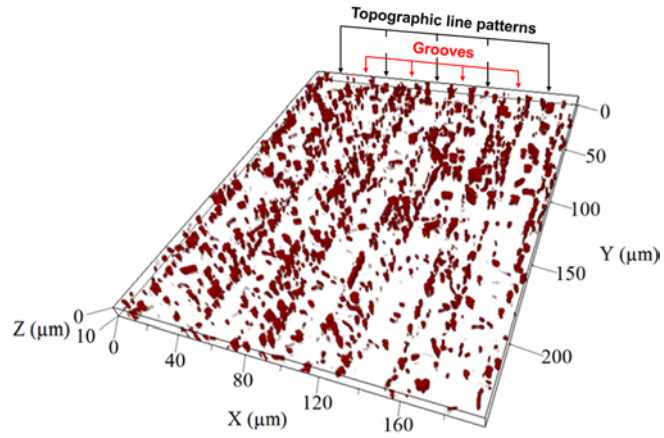


Figure S1. Representative 3D fluorescence image of WT *E. coli* biofilm. The image shows a PDMS surface modified with 20 μm wide topographic line patterns with 20 μm inter-pattern distance. The cells on top of protruding line patterns (5 μm tall) are the focus of this study.

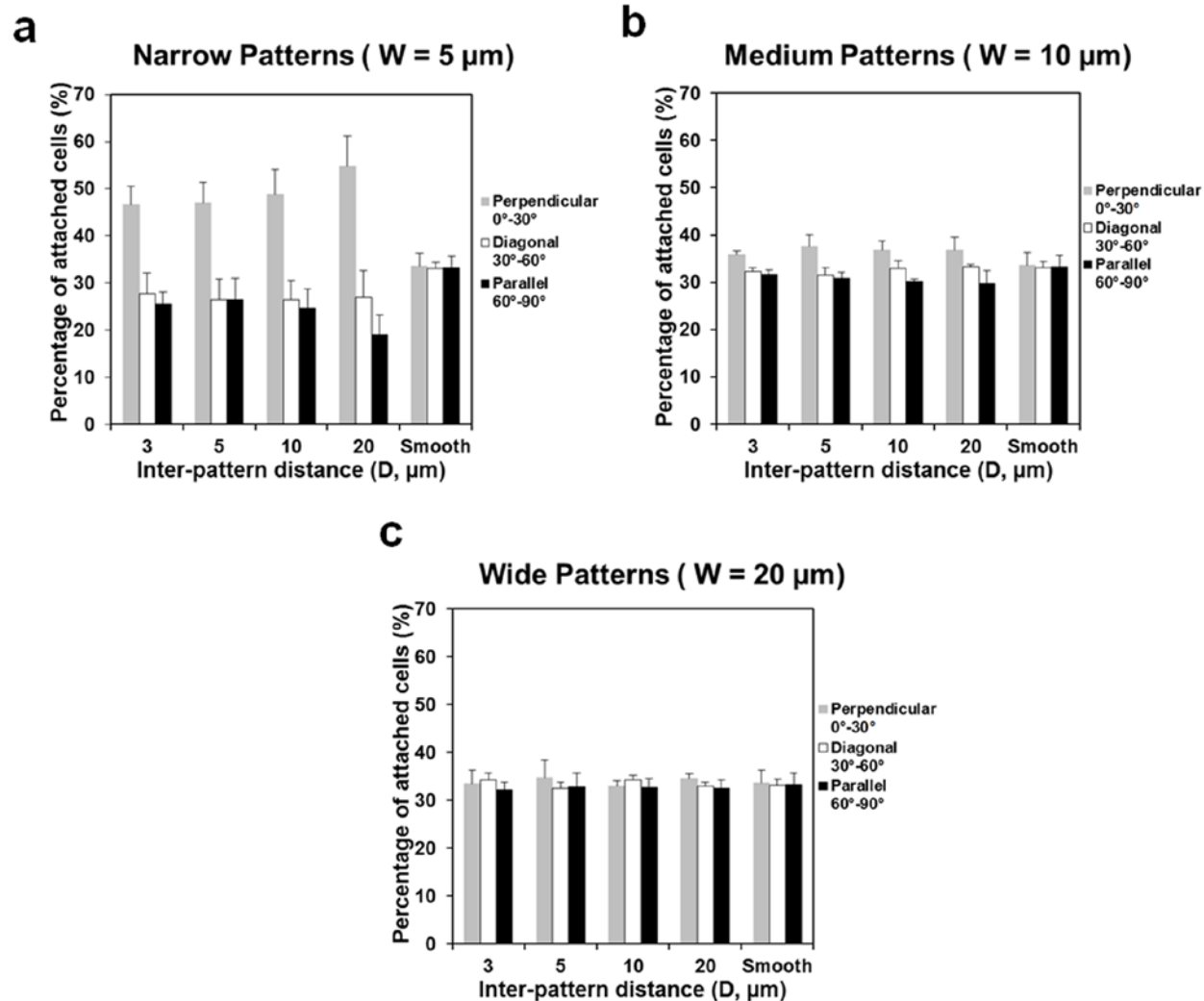


Figure S2. Orientation of the WT *E. coli* cells attached on top of PDMS line patterns at 24 h after inoculation (mean \pm standard error shown; at least three biological replicates were tested for each sample). Distribution of cell orientation on top of (a) narrow (5 μm wide), (b) medium (10 μm wide), and (c) wide (20 μm wide) PDMS line patterns.

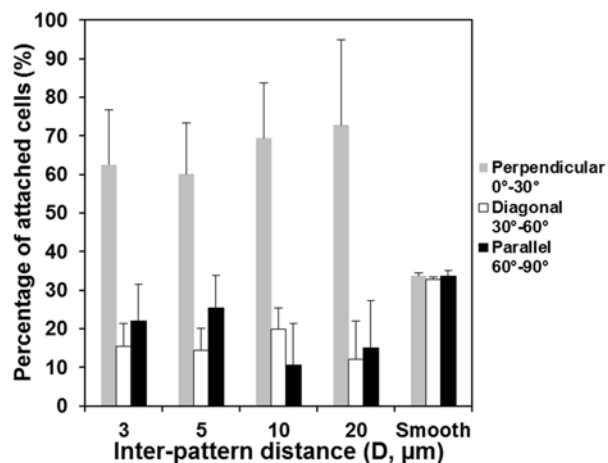


Figure S3. Orientation of WT *E. coli* cells on top of 5 μm wide line patterns and smooth PDMS surfaces after 2 h of attachment (mean \pm standard error shown; at least three biological replicates were tested for each sample).

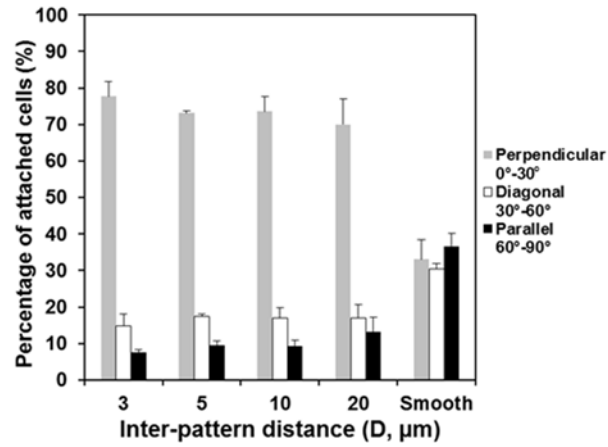


Figure S4. Orientation of WT *E. coli* cells on top of face-down 5 μm wide line patterns and smooth PDMS surfaces at 24 h after inoculation (mean \pm standard error shown; at least three biological replicates were tested for each sample).

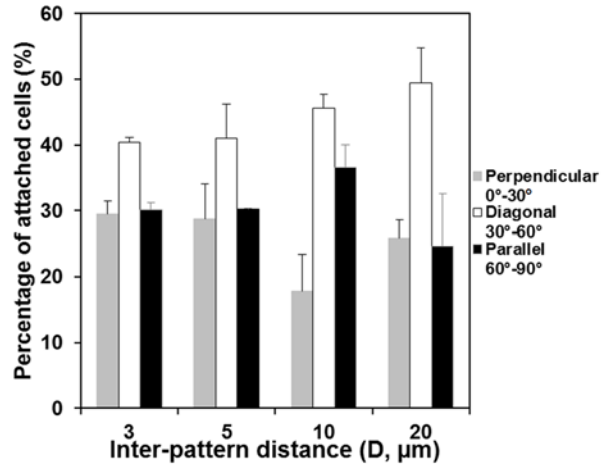


Figure S5. Orientation of WT *E. coli* cells on top of 5 μm wide line patterns after 24 h of culturing (mean ± standard error shown; at least three biological replicates were tested for each sample). Seeding cells were pretreated with 10 μg/mL chloramphenicol for 1 h before inoculation.

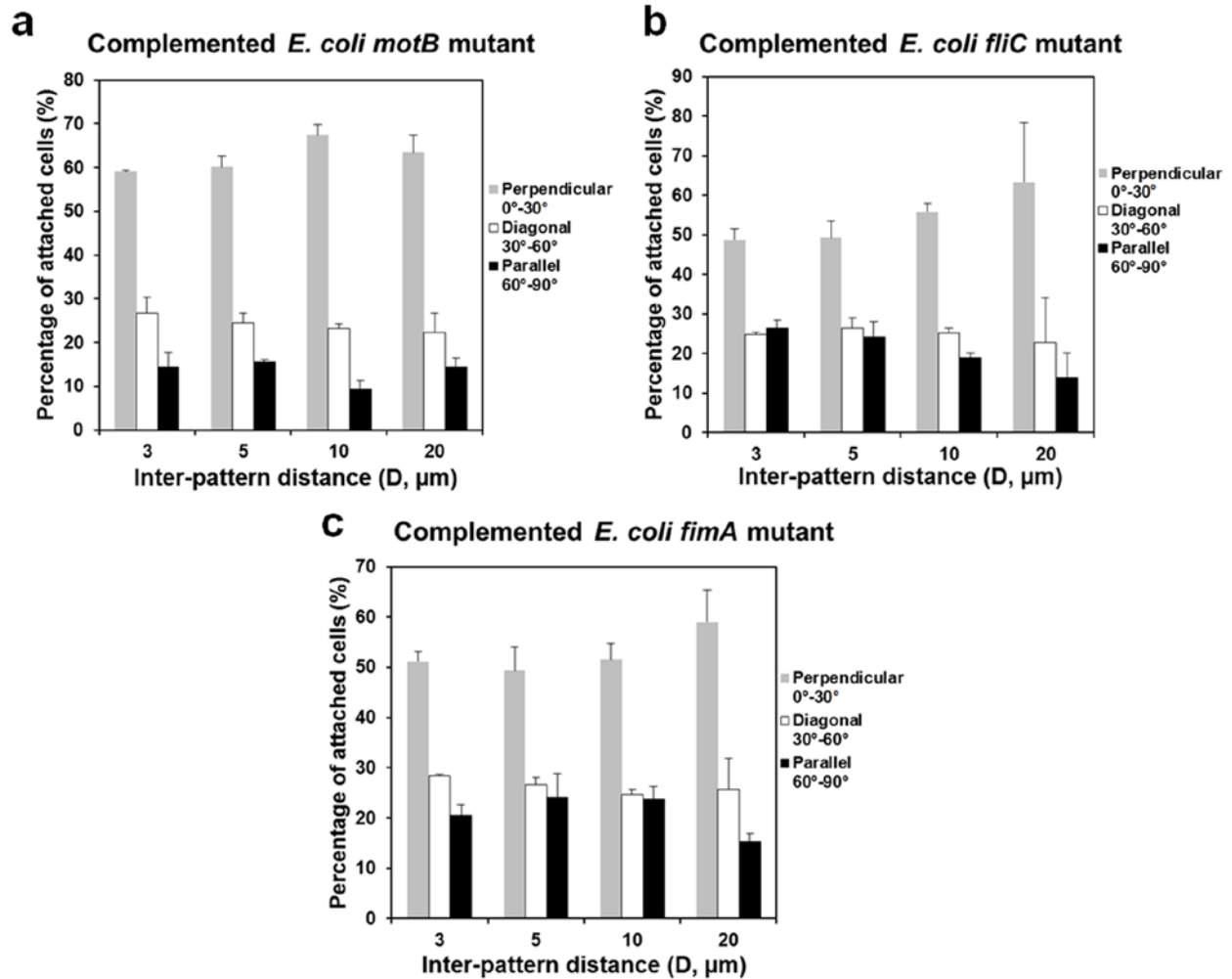


Figure S6. Distribution of cell orientation on top of 5 μm wide line patterns (mean \pm standard error shown; at least three biological replicates were tested for each sample). (a) The complemented *motB* mutant *E. coli* RP3087/pRHG03. (b) The complemented *fliC* mutant *E. coli* RHG01/pRHG04. (c) The complemented *fimA* mutant *E. coli* RHG02/pRHG05.

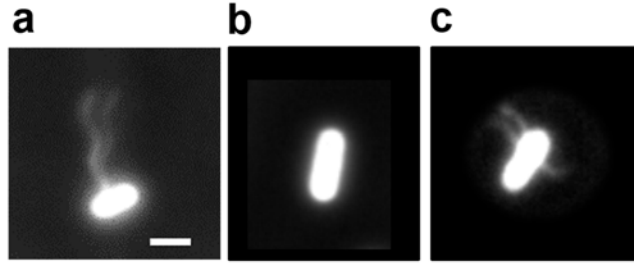


Figure S7. Fluorescence images of the planktonic cells showing flagella-like structures. The images show the cells of the WT *E. coli* (a) and its mutants of *fliC* (b) and *fimA* (c) genes. The cells harvested from planktonic cultures ($OD_{600} = 0.6$) were stained with Alexa Fluor® 594 NHS Ester (Succinimidyl Ester) Bar = 2 μm .

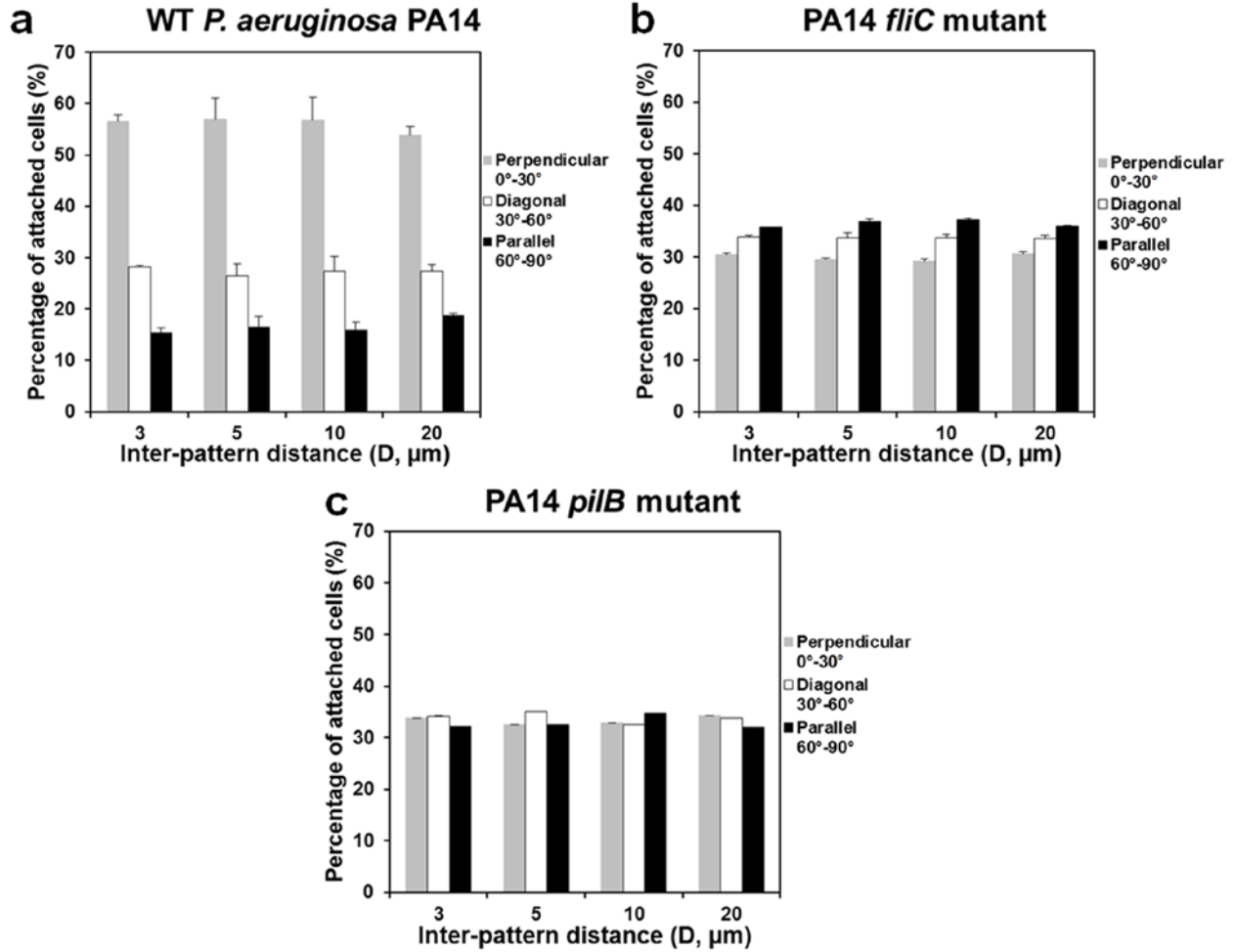


Figure S8. Distribution of cell orientations on top of 5 μm wide line patterns (mean \pm standard error shown; at least three biological replicates were tested for each sample). **(a)** Wild-type *P. aeruginosa* strain PA14. **(b)** $\Delta fliC$ mutant of *P. aeruginosa* PA14 (PA1092). **(c)** $\Delta pilB$ mutant of *P. aeruginosa* PA14 (PA4526).

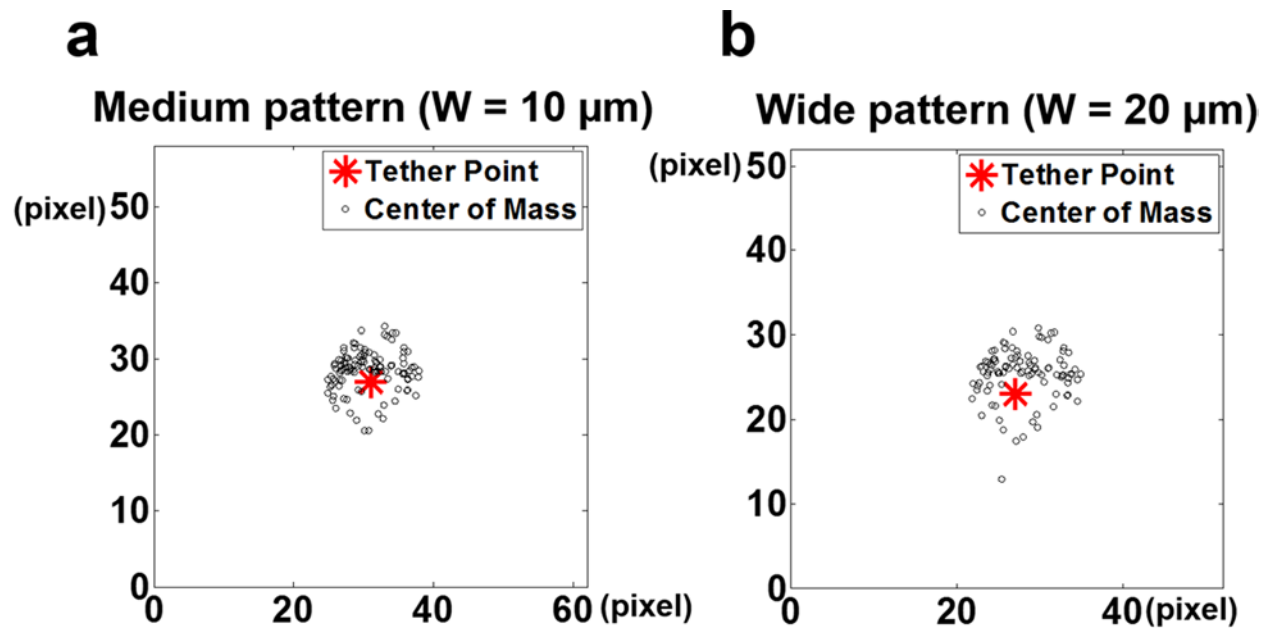


Figure S9. Rotation trajectories of cells tethered on top of medium (a) and wide (b) line patterns. The unit of x and y-axis is $0.16 \mu\text{m}/\text{pixel}$. The images were taken with 1.6 s interval between every two frames.

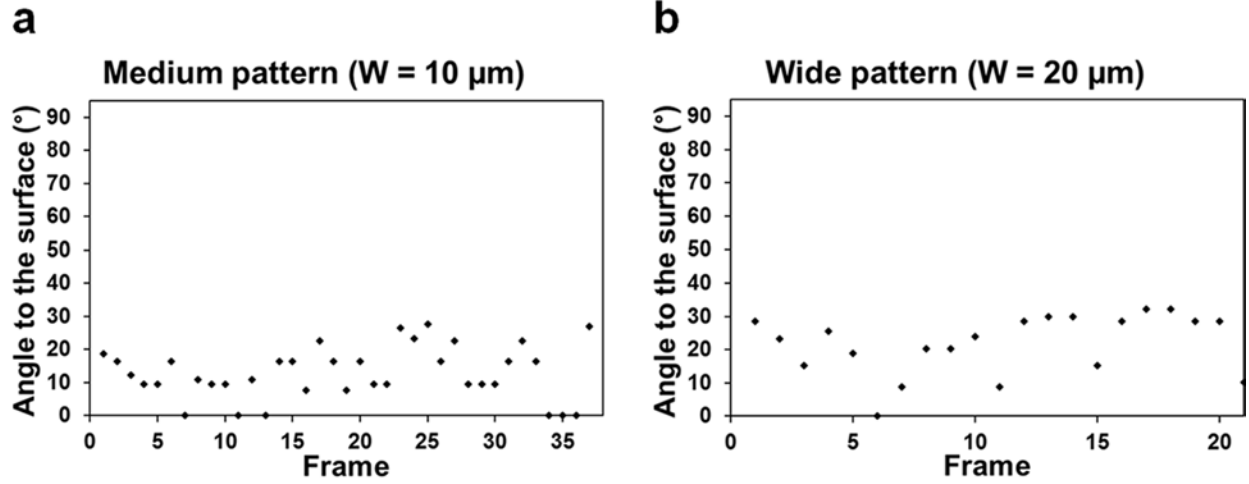


Figure S10. The angle between cell body and the substratum surface (x-y plane) of WT *E. coli* cells tethered on the top of medium (a) and wide (b) line patterns. The images were taken with 1.6 s interval between every two frames.

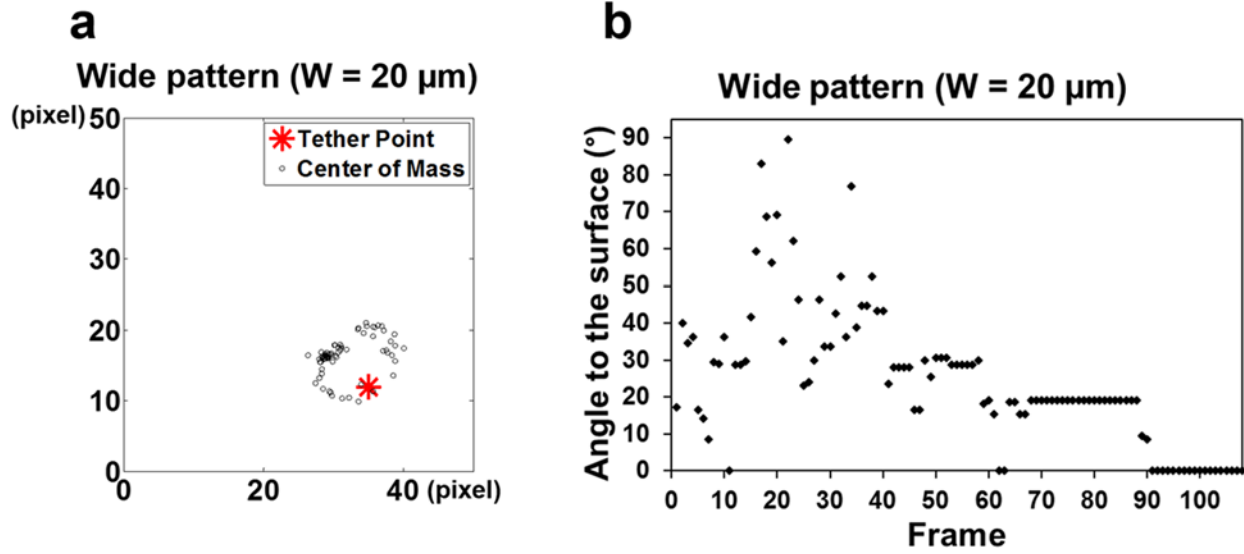


Figure S11. Rotation trajectory (a; unit of x and y-axis is $0.16 \mu\text{m}/\text{pixel}$) and the angle between cell body and the substratum surface (x-y plane) (b) of a cell tethered on the vertical side of a wide line pattern. The cell rotation pattern was still a circle but was not centered around the tether point. The images were taken with 1.6 s interval between every two frames.

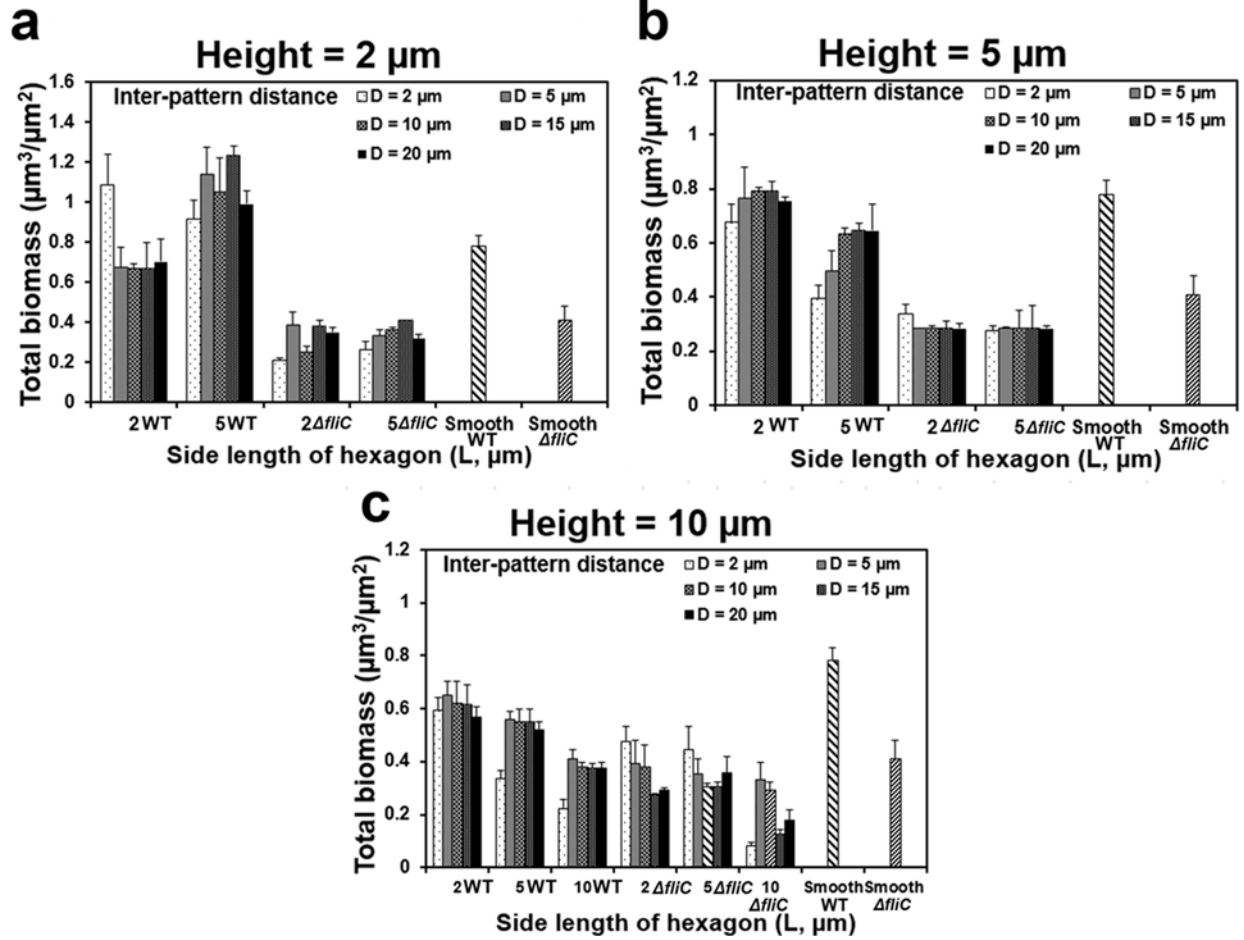


Figure S12. Biofilm formation of the WT *E. coli* and its isogenic *fliC* mutant on PDMS surfaces modified with hexagon shaped topographic patterns (mean \pm standard error shown; at least three biological replicates were tested for each sample). The graphs show the biomass of biofilms on surfaces modified with 2 (a), 5 (b), or 10 (c) μm tall hexagon shaped patterns with varied side length (2, 5, or 10 μm) and inter-pattern distance (2, 5, 10, 15, or 20 μm).

Movie S1. Rotation of a WT *E. coli* cell tethered on top of a 20 μm wide protruding line. The cell shape (circled in red) was automatically detected by ACTIVE. The movie shows that the center of cell mass followed a regular circle during rotation (based on images collected over 198 s). A few other cells were also seen with short presence during the movie. These are planktonic cells and not the focus of this study.

Movie S2. Rotation of WT *E. coli* cells on the edge of a narrow (5 μm wide) protruding line. The movie shows that the cell body moved in an irregular pattern (based on images collected over 41 s).

References:

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