Supporting Information

A particle self-aligning, focusing, and electric impedance micro-cytometer device for label-free single cell morphology discrimination and yeast budding analysis Xinwu Xie^{a,c§}, Zhiwei Zhang^{a,b§}, Xiang Ge^{a,b}, Xiaohao Zhao^{a,b}, Limei Hao^{a,c}, Zhen Cheng^d, Weibin Zhou^b, Yaohua Du^{a,c}, Lei Wang^e, Feng Tian^{a*}, Xinxi

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Figure S1. Scatter plots of beads and rods. a, scatter plot of R and amplitude; b, scatter plot of R and width. The beads-5, beads-8 and rods-8 represent \emptyset 5, 8 μ m-beads and 8 μ m-rods, respectively.



Figure S2. Histograms of ~40 typical budding yeast's orientation angle before the constriction part (a) and at the sensing zone (b).



Figure S3. Histogram of pulse width (a), amplitude (b), and R (c) of typical single yeast and budding yeast with a high speed manner. Inserts highlight the ROC curves of each parameter.



Figure S4. Typical images of yeast under different condition (p0 and p12 h represent the dry yeast powder resuscitation for 0 and 12 h respectively; 0, 12, 24, and 48 h represent fully recovered yeast cultured in fresh medium for 0, 12, 24, and 48 h, respectively).



Figure S5. The correlation coefficient of the late-budding rate data



Figure S6. The late budding rate versus the OD value.