

# A dynamic microfluidic culture model of the female reproductive tract

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The Fallopian tube epithelium plays a critical role in human fertility by facilitating gamete transport, fertilization, and early embryo development<sup>1</sup>. The cooperative activity of ciliated and secretory epithelial cells is essential for maintaining luminal homeostasis<sup>2</sup>. This balance is dynamically regulated by both mechanical and hormonal cues<sup>3</sup>. However, the role of the dynamic tubal flow environment in regulating epithelial cell behaviour, and the mechanotransduction pathways underlying their responses, remain largely undefined. Here, we present a microfluidic model that mimics the native flow microenvironment of the Fallopian tube to examine how physiological shear regulates epithelial differentiation. Primary epithelial cells were cultured in a dual-channel PDMS device under static and dynamic flow conditions mimicking *in vivo* shear rates. Cellular responses were evaluated for ciliogenesis (acetylated  $\alpha$ -tubulin, FOXJ1) and secretory activity (OVGP1). Our findings demonstrate that fluid shear promotes cilia formation by up to 3-fold and enhances OVGP1 secretion, suggesting shear-induced functional maturation of both ciliated and secretory epithelial cells. This dynamic microfluidic culture model offers a valuable platform for advancing our understanding of reproductive tract biology.

## References:

<sup>1</sup> Dinh, H.Q.; Lin, X.; Abbasi, F.; et al. Cell Rep. **2021**, 35(2), 108978.

<sup>2</sup> Abdul Halim, M.S.; Dyson, J. M.; Gong, M.; et al. Nat. Commun. **2024**, 15(1), 7411.

<sup>3</sup> Weigert, M.; Li, Y.; Zhu, L.; et al. Nat. Commun. **2025**, 16(1), 372.