Abenza et al, Supplementary Material Figure S2

The diagram shows a scatter plot with two axes: penetration (%) on the y-axis and average curvature of curved cells (µm⁻¹) on the x-axis. Several strains are plotted, including wild type, mto2Δ, tea2Δ, tea1Δ, swc2Δ, ma3Δ, alp14Δ, tea4Δ, tip1Δ, mto1Δ, vps71Δ, pom1Δ, rpi3702Δ, ria1Δ, mss116Δ, and mgr2Δ. The data points are connected with error bars to represent the variability of the measurements.