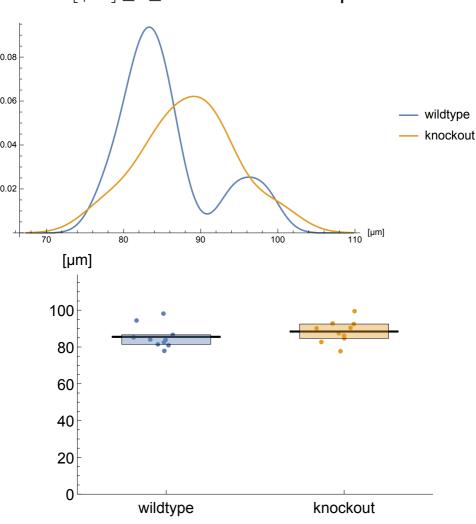
## file\_level - 7\_perimeterLength $\label{eq:main_perimeter} \begin{subarray}{c} [\mu\textbf{m}] $\_0$_cells minimal per file \end{subarray}$



genotyp	# files	# cells	7_perimeterLength [µm] mean
knockout	10	243	88.408
wildtype	10	112	85.515
total	20	355	86.9615

compared groups	statistic			
		Statistic	P-Value	
	Mann-Whitney	67.	0.212294	
	Paired T	1.09871	0.300418	
knockout/wildtype	Paired Z	1.09871	0.271897	
Knockout/ Whatype	Sign	7	0.34375	
	Signed-Rank	37.	0.358951	
	T	1.05112	0.307111	
	Z	1.05112	0.293203	