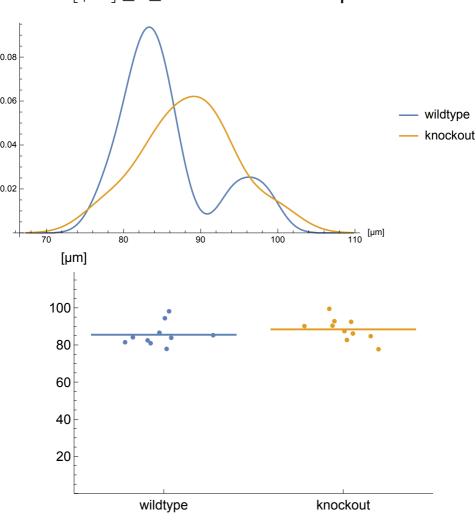
file_level - 7_perimeterLength $\label{eq:main_perimeter} \begin{subarray}{c} [\mu\textbf{m}] $_0$_cells minimal per file \end{subarray}$



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