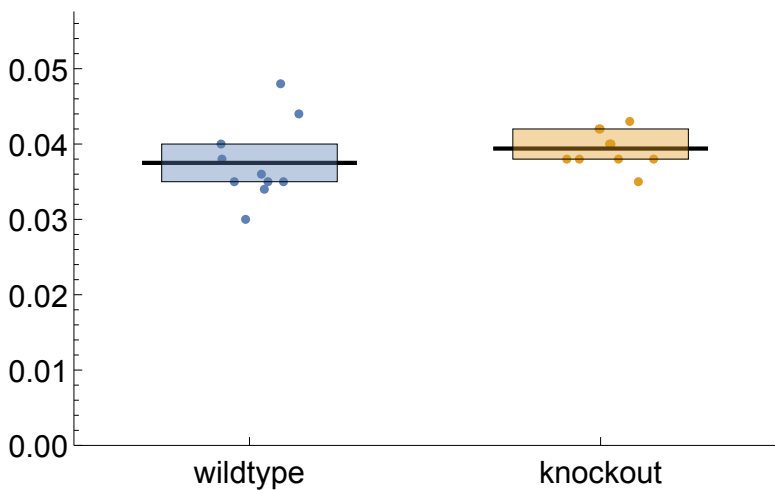
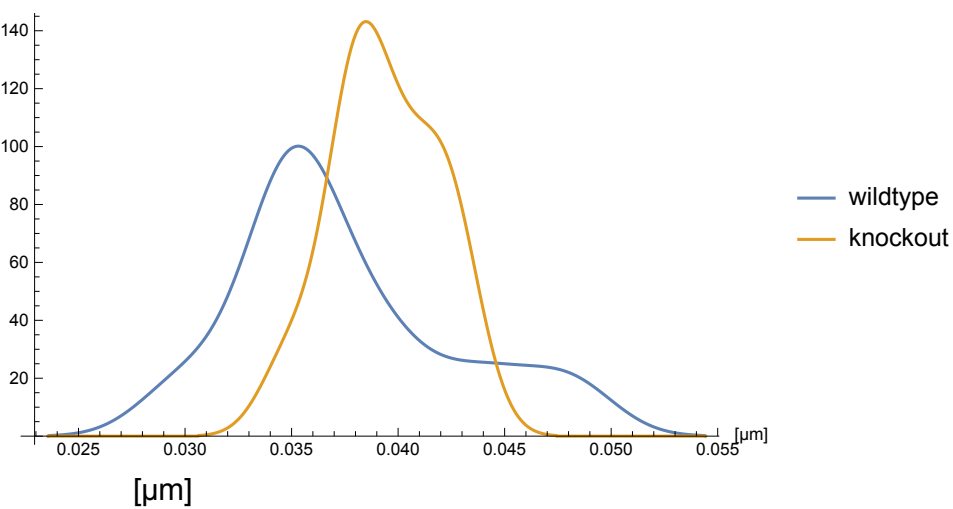


file_level - 10_minCentroidDistance
[μm]_0_cells minimal per file



genotyp	# files	# cells	10_minCentroidDistance [μm] mean
knockout	10	243	0.0394
wildtype	10	112	0.0375
total	20	355	0.03845

compared groups	statistic		
knockout / wildtype		Statistic	P-Value
	Mann-Whitney	68.5	0.167914
	Paired T	1.27808	0.233199
	Paired Z	1.27808	0.201222
	Sign	6	0.507813
	Signed-Rank	32.5	0.259555
	T	1.03568	0.314066
	Z	1.03568	0.300352