Fixed effects	Model 2a output		
	Estimate	SE	<i>P</i> -value <sup>1</sup>
Intercept <sup>a</sup>	-4.853 x 10 <sup>-1</sup>	1.593 x 10 <sup>-1</sup>	0.002
penicillins	-1.693 x 10 <sup>-1</sup>	6.472 x 10 <sup>-2</sup>	0.009
cloxacillin	-2.334 x 10 <sup>-1</sup>	3.882 x 10 <sup>-2</sup>	< 0.001
cephalosporins	-3.212 x 10 <sup>-1</sup>	9.738 x 10 <sup>-2</sup>	0.001
rifaximin	-1.105 x 10 <sup>-1</sup>	1.800 x 10 <sup>-1</sup>	0.539
Parity	-3.112 x 10 <sup>-2</sup>	9.907 x 10 <sup>-3</sup>	0.002
sqrt(milk yield before) <sup>b</sup>	3.080 x 10 <sup>-3</sup>	1.778 x 10 <sup>-3</sup>	0.084
Random effects	Variance	SD	
Animal	2.320 x 10 <sup>-1</sup>	4.816 x 10 <sup>-1</sup>	
Herd	1.209 x 10 <sup>-2</sup>	1.100 x 10 <sup>-1</sup>	
residuals	1.563 x 10 <sup>-1</sup>	3.953 x 10 <sup>-1</sup>	

Supplement Table S3. Model 2a: Results from general linear mixed model testing the association between cell count ratio and different antibiotic dry cow therapy groups in 1,380 cows from 237 dairy farms

<sup>1</sup>Significance was declared at *P*-values <0.05.

<sup>a</sup>Intercept = cell count ratio was the reference category and calculated using the SCC from the last milk recording data before dry cow therapy and the first SCC after calving. <sup>b</sup>sqrt(milk yield before) = square root from the milk yield in 305 DIM from the milk recording data before dry cow therapy.