

Species	N positive	Prevalence (%)	95% CI (%)
<i>Cooperia oncophora</i>	8	88.9	56.5 – 98.0
<i>Cooperia punctata</i>	5	55.6	26.7 – 81.1
<i>Cooperia</i> spp.	5	55.6	26.7 – 81.1
<i>Ostertagia ostertagi</i>	9	100	70.1 - 100
<i>Ostertagia leptospicularis</i>	2	22.2	6.3 – 54.7
<i>Oesophagostomum radiatum</i>	4	44.4	18.9 – 73.3
<i>Oesophagostomum venulosum</i>	4	44.4	18.9 – 73.3
<i>Trichostrongylus axei</i>	4	44.4	18.9 – 73.3
<i>Trichostrongylus colubriformis</i>	1	11.1	2.0 – 43.5
<i>Trichostrongylus</i> spp.	2	22.2	6.3 – 54.7
<i>Chabertia ovina</i>	1	11.1	2.0 – 43.5
<i>Bunostomum phlebotomum</i>	3	33.3	12.1 – 64.6
<i>Haemonchus contortus</i>	1	11.1	2.0 – 43.5
Strongylidae	1	11.1	2.0 – 43.5

Table S2 Farm prevalence with 95% confidence intervals (CI) of strongyle species identified by deep amplicon sequences in nine German cattle farms before treatment.