

# Supplementary Materials: Aflatoxin B1 Impairs Bone Mineralization in Broiler Chickens

**Supplementary Table S1.** Effect of increasing dosage of Aflatoxin B1 contaminated diet on microstructure of femoral metaphysis on day 8 <sup>1</sup>.

Bone Region	Parameters <sup>2</sup>	Unit	Treatment Groups <sup>3</sup>					SEM	P-values <sup>4</sup>		
			T1	T2	T3	T4	T5		P <sub>Model</sub>	P <sub>Linear</sub>	P <sub>Quadratic</sub>
Total	BMC	g	0.016	0.019	0.018	0.018	0.020	0.001	0.5944	0.2700	0.81380
	BMD	g/mm <sup>3</sup>	0.28	0.33	0.32	0.31	0.34	0.02	0.1163	0.0907	0.4325
	TV	mm <sup>3</sup>	58.2	56.2	56.5	56.7	57.9	2.8	0.9817	0.9934	0.5577
	BV	mm <sup>3</sup>	25.8	29.9	29.0	27.4	31.3	2.5	0.5777	0.2999	0.8889
	BV/TV	-	44.2	52.6	51.2	47.7	53.9	2.7	0.1104	0.1042	0.5284
	TS	mm <sup>2</sup>	94	92	94	93	94	3	0.9887	0.9255	0.6830
	BS	mm <sup>2</sup>	165	155	159	163	153	10	0.8961	0.6146	0.9804
	BS/BV	mm <sup>-1</sup>	6.6	5.4	5.6	6.1	5.0	0.4	0.0619	0.0484	0.6849
	BS/TV	mm <sup>-1</sup>	2.84	2.77	2.81	2.86	2.65	0.09	0.4817	0.3110	0.4111
Cortical	BMC	g	0.014	0.017	0.016	0.015	0.017	0.001	0.3981	0.2452	0.6860
	BMD	g/mm <sup>3</sup>	0.506	0.544	0.531	0.530	0.538	0.013	0.3768	0.2750	0.3784
	TV	mm <sup>3</sup>	26.6	30.9	30.2	28.2	32.1	2.6	0.6387	0.3439	0.8409
	BV	mm <sup>3</sup>	25.2	29.5	28.7	26.8	30.9	2.4	0.5313	0.2820	0.8853
	BV/TV	-	94.8	95.8	95.1	95.3	96.5	0.7	0.5683	0.2634	0.6404
	TS	mm <sup>2</sup>	135.2	129.3	131.2	132.4	131.3	3.8	0.8718	0.7110	0.5491
	BS	mm <sup>2</sup>	154.7	147.5	152.5	153.4	147.4	8.5	0.9614	0.7562	0.9571
	BS/BV	mm <sup>-1</sup>	6.36	5.19	5.40	5.84	4.82	0.38	0.0850	0.0616	0.7310
	BS/TV	mm <sup>-1</sup>	6.03	4.97	5.13	5.57	4.64	0.35	0.0954	0.0700	0.6872
Trabecular	BMC	g	0.0021	0.0018	0.0017	0.0020	0.0017	0.0001	0.3461	0.3300	0.4285
	BMD	g/mm <sup>3</sup>	0.0736	0.0810	0.0745	0.0770	0.0761	0.0036	0.6877	0.9246	0.5888
	TV	mm <sup>3</sup>	28.12	21.96	23.17	25.20	22.76	1.42	0.0621	0.1334	0.1597

<b>BV</b>	mm <sup>3</sup>	0.222	0.181	0.144	0.244	0.128	0.048	0.4560	0.4510	0.9460
<b>BV/TV</b>	-	0.786	0.847	0.623	0.927	0.509	0.181	0.5473	0.4436	0.5545
<b>TS</b>	mm <sup>2</sup>	63.2	53.6	55.5	58.6	54.4	2.1	0.0373	0.0898	0.1657
<b>BS</b>	mm <sup>2</sup>	10.5	7.4	6.6	10.0	5.6	1.7	0.2382	0.2152	0.8169
<b>BS/BV</b>	mm <sup>-1</sup>	50.0	45.7	46.7	44.7	48.3	3.4	0.8270	0.6861	0.3298
<b>BS/TV</b>	mm <sup>-1</sup>	0.379	0.346	0.287	0.385	0.227	0.061	0.3690	0.2021	0.6971

<sup>1</sup> Data represents mean values of six femur bone samples per treatment. SEM represents the pooled standard error of the mean for each parameter across all treatment groups.

<sup>2</sup> Parameters: BMC, bone mineral content; BMD, bone mineral density; TV, tissue volume; BV, bone volume; BV/TV, bone volume/ tissue volume; TS, tissue surface area; BS, bone surface area; BS/BV, bone surface area/ bone volume; BS/ TV, bone surface area/ tissue volume.

<sup>3</sup> Treatment groups: T1 (< 2 ppb AFB1), T2 (75-79 ppb AFB1), T3 (150 ppb AFB1), T4 (230-260 ppb AFB1), T5 (520-560 ppb AFB1).

<sup>4</sup> P-values: P<sub>Model</sub>, p-value for the whole model; P<sub>Linear</sub>, p-value for the linear regression; P<sub>Quadratic</sub>, p-value for the quadratic regression

**Supplementary Table S2.** Effect of increasing dosage of Aflatoxin B1 contaminated diet on tibial bone parameters on day 8 <sup>1</sup>.

Parameters <sup>2</sup>	Treatment Groups <sup>3</sup>					SEM	P-values <sup>4</sup>		
	T1	T2	T3	T4	T5		P <sub>Model</sub>	P <sub>Linear</sub>	P <sub>Quadratic</sub>
FBW (g)	0.983	0.982	1.031	1.018	1.053	0.055	0.8879	0.3466	0.96515
DBW (g)	0.327	0.329	0.339	0.342	0.353	0.018	0.8516	0.2704	0.8540
FFDW (g)	0.322	0.330	0.335	0.333	0.356	0.020	0.8419	0.3077	0.7721
AW (g)	0.1454	0.1525	0.1548	0.1512	0.1656	0.0108	0.7745	0.2750	0.8342
Ash %	44.92	46.11	46.06	45.38	46.47	0.64	0.5247	0.2892	0.7509

<sup>1</sup> Data represents mean values of six tibia bone samples per treatment. SEM represents the pooled standard error of the mean for each parameter across all treatment groups.

<sup>2</sup> Parameters: FBW, fresh bone weight; DBW, dry bone weight; FFDW, fat free dry weight; AW, ash weight.

<sup>3</sup> Treatment groups: T1 (< 2 ppb AFB1), T2 (75-79 ppb AFB1), T3 (150 ppb AFB1), T4 (230-260 ppb AFB1), T5 (520-560 ppb AFB1).

<sup>4</sup> P<sub>Model</sub>, p-value for the whole model; P<sub>Linear</sub>, p-value for the linear regression; P<sub>Quadratic</sub>, p-value for the quadratic regression.

**Supplementary Table S3.** Effect of increasing dosage of Aflatoxin B1 contaminated diet on mRNA levels of calcium and phosphorus transporter genes on day 8 <sup>1</sup>.

Genes <sup>1</sup>	Treatment Groups <sup>3</sup>					SEM	P-values <sup>4</sup>		
	T1	T2	T3	T4	T5		P <sub>Model</sub>	P <sub>Linear</sub>	P <sub>Quadratic</sub>
CALB1	1.00	1.07	1.80	1.62	0.87	0.39	0.4116	0.8166	0.1109
CASR	1.00	0.71	0.75	0.68	0.63	0.22	0.7679	0.2665	0.6418
NCX1	1.00	0.66	0.72	0.76	1.02	0.23	0.6989	0.8495	0.1713
NaPi-IIb	1.00	1.25	1.20	1.53	1.10	0.21	0.4980	0.4868	0.2370
PMCA1	1.00 <sup>ab</sup>	1.40 <sup>a</sup>	0.94 <sup>ab</sup>	0.48 <sup>ab</sup>	0.33 <sup>b</sup>	0.25	0.0395	0.0086	0.2407
VDR	1.00	1.89	2.16	1.41	2.26	0.38	0.1530	0.1044	0.4585

<sup>a-b</sup> Means within a column not sharing a common letter differ significantly ( $P < 0.05$ ).

<sup>1</sup> Data are presented as mean values of fold change in mRNA expression of calcium and phosphorus transporter genes using  $2^{-\Delta\Delta C_t}$  method. Data relates to 6 jejunum samples per treatment.

<sup>2</sup> Genes: CALB1, calbindin 1; CaSR, calcium-sensing receptor; NCX1, sodium-calcium exchanger 1; NaPi-IIb, sodium-phosphate cotransporter; PMCA1b, plasma membrane calcium ATPase 1b; VDR, vitamin D receptor.

<sup>3</sup> Treatment groups: T1 (< 2 ppb AFB1), T2 (75-79 ppb AFB1), T3 (150 ppb AFB1), T4 (230-260 ppb AFB1), T5 (520-560 ppb AFB1).

<sup>4</sup> P<sub>Model</sub>, p-value for the whole model; P<sub>Linear</sub>, p-value for the linear regression; P<sub>Quadratic</sub>, p-value for the quadratic regression.

**Supplementary Table S4.** Effect of increasing dosage of Aflatoxin B1 contaminated diet on mRNA levels of tight junction proteins on day 8 <sup>1</sup>.

Genes <sup>2</sup>	Treatment Groups <sup>3</sup>					SEM	P-values <sup>4</sup>		
	T1	T2	T3	T4	T5		P <sub>Model</sub>	P <sub>Linear</sub>	P <sub>Quadratic</sub>
MUC2	1.00	0.91	1.14	0.80	1.28	0.20	0.6294	0.5558	0.5192
CLDN1	1.00	0.48	1.21	0.57	0.95	0.26	0.3939	0.9944	0.6982
ZO1	1.00	0.72	0.85	0.57	0.75	0.12	0.1842	0.1092	0.284
OCLN	1.00	1.00	1.09	1.06	0.99	0.11	0.9715	0.9103	0.5989

<sup>1</sup> Data are presented as mean values of fold change in mRNA expression of tight junction related genes using  $2^{-\Delta\Delta Ct}$  method. Data relates to 6 jejunum samples per treatment.

<sup>2</sup> Genes: MUC2, mucin 2; CLDN1, claudin 1; ZO1, zonula occludens; and OCLN, occluding.

<sup>3</sup> Treatment groups: T1 (< 2 ppb AFB1), T2 (75-79 ppb AFB1), T3 (150 ppb AFB1), T4 (230-260 ppb AFB1), T5 (520-560 ppb AFB1).

<sup>4</sup> P<sub>Model</sub>, p-value for the whole model; P<sub>Linear</sub>, p-value for the linear regression; P<sub>Quadratic</sub>, p-value for the quadratic regression.