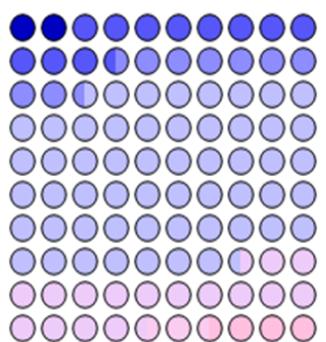
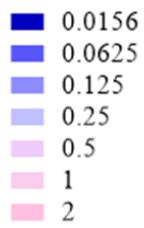


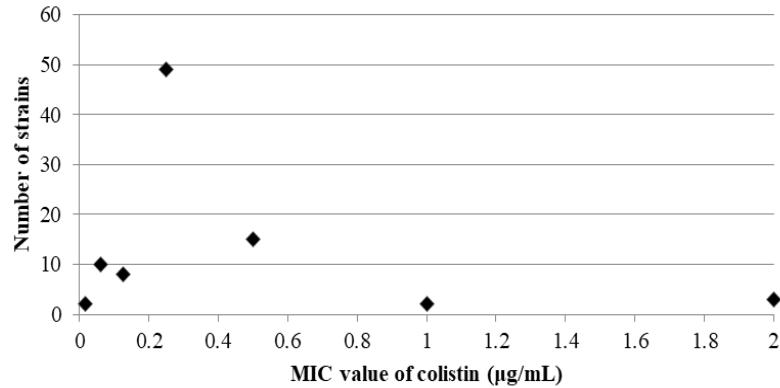
Figure S1. Identification of avian pathogenic *Escherichia coli* (APEC) strains by PCR. Lane (a) shows the expression of *fimAvMT78*, (b) is *tsh*, (c) is *felA*, (d) is *fimH*, (e) is *sta*, (f) is *yaiO*, (g) is *fimA*, (h) is *papC*, (i) is *iutA*, (j) is *papGIA2*, and (k) is *papGJ96*. Primer information is in Table S1. Horizontal numbers (1–20) mean each APEC strain, E means *E. coli* ATCC 25922, and S means *Salmonella enterica* subsp. *enterica* serovar Typhimurium ATCC 14028 and N means Negative control.



Total=89



MIC distribution of isolated strains



a

b

Figure S2. MIC distribution of isolated APECs. MICs of 89 APECs were confirmed less than $2 \mu\text{g/mL}$, that means colistin susceptible [49] (a). 49 APECs has $0.25 \mu\text{g/mL}$ MIC value, and it was determined MIC₅₀ (b).

Table S1. Primer information.

Primer *	Primer Sequence	Size	Tm (°C)	Accession Number	Reference
APEC_fimAv _{MT78} _F	tctggctgatactacacc	266	52	Z37500	[38]
APEC_fimAv _{MT78} _R	actttagatgagtagctg				
APEC_tsh_F	ggtgtgcactggagtgg	640	58	L27423	[39]
APEC_tsh_R	agtccagcgtatagttgg				
APEC_felA_F	ggtcaascagctaaaaacggtaagg	239	61	GCA_001620375.1	[40]
APEC_felA_R	ccttcagaaacagttaccgcattcg				
APEC_fimH_F	gatcttcgacgcaatc	389	52	AJ225176.1	[41]
APEC_fimH_R	cgagcagaaacatcgcatcg				
APEC_sta_F	ttaatagcacccggtaacaaggcagg	147	55	GPL3935	[32]
APEC_sta_R	cttgacttccaaaagagaaaattac				
E. coli_yaiO_F	tgatttccgtgcgtctgaatg	115	58	EG13297	[42]
E. coli_yaiO_R	atgctccgttagcggttttc				
APEC_fimA_F	cggtctgtccctsagt	500	52	AF490890	[43]
APEC_fimA_R	gtcgcatccgcattagc				
APEC_papC_F	gacggctgtactgcaggggtgtggcg	328	61	DQ010312.1	[44]
APEC_papC_R	atatccttctgcagggatgcaata				
APEC_iutA_F	atgagcatatctccggacg	587	58	GCA_001021615.1	[33]
APEC_iutA_R	caggctcgagaacatctgg				
APEC_papGIA2_F	gggatgagcccccccttgat	190	63		[45]
APEC_papGIA2_R	cggggccccaagtaactcg				
APEC_papGJ96_F	tcgtgtcaggcggatatt	461	63		[46]
APEC_papGJ96_R	tggcatcccccaacattatcg				

* The primers are designed to target the following adhesins: type 1 pili (*fim*), F11 fimbriae (*felA*), aerobactin presence (*iutA*), pili associated with pyelonephritis (*pap*), heat stable enterotoxins (*sta*), temperature-sensitive hemagglutinin (*tsh*).

Table S2. Primers used in qRT-PCR gene expression.

Target gene	Sequence (5'-3')
<i>pmrA</i>	F: acggctgccttatgcctac R: tagcagataaatagcgtcagg
<i>pmrB</i>	F: atatccataaccctgcgcgac R: gatgatattgaccacgagatt
<i>mgrB</i>	F: ttcatcacgcctgaattac R: cgactcattccgaaaaagca
<i>phoP</i>	F: gggttaactcgacatgaaact R: ttcaacttacccctcc
<i>phoQ</i>	F: aaagccgcgtttaacacca R: atgtactgtggacgtctg
<i>mcr-1</i>	F: cggaaattccaagatacaaattataaatactct R: acgcgtcgactcagcgatgaatcggtcggtc

Table S3. In vitro pharmacodynamic profile of isolates

	APEC 1	APEC 2	APEC 3	APEC 4	APEC 5	APEC 6	APEC 7	APEC 8	APEC 9	APEC 10
MIC	0.03–0.06	0.06–0.13	0.016	0.06–0.13	0.06	0.25	0.02–0.06	0.25–0.50	0.03–0.06	0.06–0.25
MBC	0.03–0.06	0.06–0.13	0.016	0.06–0.13	0.06	0.25	0.02–0.06	0.25–0.50	0.03–0.06	0.06–0.25
MBC/MIC	1	1	1	1	1	1	1	1	1	1
	APEC 11	APEC 12	APEC 13	APEC 14	APEC 15	APEC 16	APEC 17	APEC 18	APEC 19	APEC 20
MIC	0.06–0.13	0.0625	0.03–0.06	0.03–0.06	0.03–0.06	0.06–0.13	0.03–0.06	0.25	0.0156	0.06–0.13
MBC	0.06–0.13	0.0625	0.03–0.06	0.0625	0.03–0.06	0.06–0.13	0.0625	0.25	0.0156	0.125
MBC/MIC	1	1	1	1	1	1	1	1	1	1
	APEC 21	APEC 22	APEC 23	APEC 24	APEC 25	APEC 26	APEC 27	APEC 28	APEC 29	APEC 30
MIC	0.13	0.25	0.25	0.25	0.50	0.25	2.00	0.25	0.25	0.25
MBC	0.13	0.25	0.25	0.50	1.00	0.25	2.00	0.25	0.25	0.25
MBC/MIC	1	1	1	2	2	1	1	1	1	1
	APEC 31	APEC 32	APEC 33	APEC 34	APEC 35	APEC 36	APEC 37	APEC 38	APEC 39	APEC 40
MIC	0.25	0.25	0.50	0.25	0.25	0.25–1.00	0.25	0.25	0.25	0.25
MBC	0.25	0.25	0.50	0.25	0.25	2	0.25	0.25	0.25	0.25
MBC/MIC	1	1	1	1	1	2	1	1	1	1
	APEC 41	APEC 42	APEC 43	APEC 44	APEC 45	APEC 46	APEC 47	APEC 48	APEC 49	APEC 50
MIC	0.25	0.25	0.25	0.25	0.25	0.25	0.13	0.50	0.50	0.25

MBC	0.25	0.25	0.25	0.25	0.25	0.25	0.13	0.50	0.50	0.25
MBC/MIC	1	1	1	1	1	1	1	1	1	1
	APEC 51	APEC 52	APEC 53	APEC 54	APEC 55	APEC 56	APEC 57	APEC 58	APEC 59	APEC 60
MIC	0.25	0.13	0.25	0.25–0.50	0.13	0.25	0.25	0.25	0.25	0.25
MBC	0.25	0.13	0.25	0.50	0.13	0.25	0.25	0.25	0.25	0.25
MBC/MIC	1	1	1	1	1	1	1	1	1	1
	APEC 61	APEC 62	APEC 63	APEC 64	APEC 65	APEC 66	APEC 67	APEC 68	APEC 69	APEC 70
MIC	0.50	0.25	0.25–1.00	0.25	0.13–0.25	0.25	0.13–0.50	0.25–0.50	0.25	0.50
MBC	0.50	0.25	0.25–1.00	0.25	0.13–0.25	0.25	0.13–0.50	0.25–0.50	0.25	0.50
MBC/MIC	1	1	1	1	1	1	1	1	1	1
	APEC 71	APEC 72	APEC 73	APEC 74	APEC 75	APEC 76	APEC 77	APEC 78	APEC 79	APEC 80
MIC	0.50	0.25	0.13–0.25	0.25	0.25	0.25	0.25–0.50	0.50–2.00	0.13–0.25	0.50
MBC	0.50	0.25	0.13–0.25	0.25	0.25	0.25	0.25–0.50	0.50–2.00	0.13–0.25	0.50
MBC/MIC	1	1	1	1	1	1	1	1	1	1
	APEC 81	APEC 82	APEC 83	APEC 84	APEC 85	APEC 86	APEC 87	APEC 88	APEC 89	
MIC	0.50–2.00	0.13–0.25	0.25–0.50	0.25	0.13–0.25	0.25	0.25	0.25–0.50	0.25	
MBC	0.50–2.00	0.13–0.25	0.25–0.50	0.25	0.25	0.25	0.25	0.25–0.50	0.25	
MBC/MIC	1	1	1	1	2	1	1	1	1	