

**Supplementary Table 2.** Geographical distribution, species, clones, and resistance mechanisms of antibiotic-resistant Gram-positive bacteria isolated from animals in Africa from 2007–2019.

Country (n) <sup>1</sup>	Year	Organism/Species (n) <sup>2</sup>	Specimen Sources (n) <sup>3</sup>	Sample size (Resistant isolates)	Clones (n) <sup>4</sup>	Resistance genes/mechanisms (n) <sup>5</sup>	Antibiotic resistance phenotype (n) <sup>6</sup>	MGEs (n) <sup>7</sup>	Reference
Angola(1)	2015	<i>E. faecium</i> (3)	Pig feces(1), Chicken feces(2)	3	ST971, ST245(2)	<i>tet</i> (L)(1), <i>tet</i> (M)(2), <i>erm</i> (B)(2)	CIP(1),TET(3),ERY(2),STR(2),NIT(2),Q/D(2)	ND	1
Egypt (15)	2019	<i>E. faecium</i> (53), <i>E. avium</i> (14), <i>E. raffinosus</i> (14), <i>E. gallinarum</i> (6), <i>E. durans</i> (8), <i>E. faecalis</i> (11)	Chicken (30), ducks(35), pigs(41)	(106)	ND	<i>CatA</i> (20), <i>CatB</i> (20), <i>fexA</i> (15), <i>fexB</i> (11), <i>vanA</i> (20), <i>vanB</i> (27), <i>vanC</i> (35)	AMP(77),CLI(106),ERY(105),TET(104),GEN(105),CIP(105),CHL(81), <i>van</i> (101)	ND	2
	2017	<i>S. aureus</i> (3), <i>S. hycus</i> (6), <i>S. intermedius</i> (3) <i>S. epidermis</i> (1) <i>S. hemolyticus</i> (1), <i>S. hominis</i> (1), <i>S. l ugdunensis</i> (3), <i>S. simulans</i> (1), <i>S. scuri</i> (4)	imported beef meat (23)	23(16)	ND	<i>mecA</i> (5), <i>gyrA</i> (12), <i>grIA</i> (10), <i>gyrB</i> (6),	AMP(6),CHL(1),CIP(8),CLI(15), ERY(6),GEN(14),MET(8),OXA(13), ,PEN(22), TET(6)	SCC <i>mec</i> (5)	3
	2017	<i>S. aureus</i> (84)	Milk(84)	84(80)	ND	<i>mecA</i> (42), <i>blaZ</i> (67)	AMX(54), SXT(66),GEN(20),CIP(12),CHL(58),PEN(70),R IF(32),AMK(14), VAN(64),STR(50),TET(44),ERY(40), AMP(80),OXA(42)	SCC <i>mec</i> (42)	4

<sup>1</sup> Total number of studies per country

<sup>2</sup> Total number of bacteria isolated

<sup>3</sup> Total number of Specimen source

<sup>4</sup> Total number of resistant clones

<sup>5</sup> Total number of resistant genes

<sup>6</sup> Number of different antibiotics any one isolate is resistant to

<sup>7</sup> Total number of mobile genetic elements: plasmids, transposons, integrons

2016	<i>S. aureus</i> (73)	Animal (73)	73(NS)	ST113(1), ST80(1)	<i>mecA</i> (14), <i>erm</i> (C)(14)	CLI(NS),CIP(NS),GEN(NS),SXT(NS),OXA(NS),E RY(NS)	SCC <i>mec</i> c(14)	5
2016	<i>S. aureus</i> (30)	raw chicken breast fillet (40), sliced luncheon meat (20), and chicken nuggets (20), Human (18)	40 (21)	ND	<i>mecA</i> (10)	DOX(31), AMX(29), OFX(10), CFP(23), CLI(21), GEN(20), APR(16), ERY(21), SXT(23), LUX(18), NAL(20), OFX(10), CIP(16).	SCC <i>mec</i> c(10)	6
2016	<i>S. aureus</i> (70)	Bovine(70)	70(41)	ND	<i>mecA</i> (2)	CRO(41),ERY(35),OXA(41),SXT(14),GEN(14), CIP(11),CLI(8),VAN(1)	SCC <i>mec</i> c(2)	7
2016	<i>S. aureus</i> (40)	Milk(30), meat(10)	40(22)	ND	<i>erm</i> (A)(18), <i>mrs</i> (A)( 4), <i>mphC</i> (6), <i>erm</i> (B) (3)	ERY(22),CLI(4),TET(24),CIP(4),CHL(5),AMX(2 6),FOX(22),SXT(1),RIF(5),GEN(4),CRO(14)	ND	8
2016	<i>S. aureus</i> (200)	Raw milk (40), Damietta Cheese (40), Kareish cheese (40), ice cream (40), and yogurt (40)	200 (106)	ND	<i>mecA</i> (106)	TET(270), NEL(78), AMX(230), CLX(314),STR(186),SXT(58), GEN(114), PEN(364), RIF(152), CHL(128), AMK(146), VAN(36)	SCC <i>mec</i> c(106)	9
2016	<i>E. faecalis</i> (6), <i>E. gallinarum</i> (2)	Fish (8)	(8)	ND	<i>VanC</i> (3), <i>vanA</i> (3)	PEN(1),VAN(1),AP(2),ERY(6),TET(8),CIP(1),A MP(2)CHL(3),RIF(4)GEN(1)	ND	10
2015	<i>S. aureus</i> (133)	cow milk samples (61), various origins (14), minced meat (6), sausage (4) and burger (7), pus (22), sputum (17), urine (1), cerebrospinal fluid (1)	133 (96)	ND	<i>mecA</i> (30)	CRO(96), TET(90), OXA(70), FOX(65), ERY(81),VAN(4),IPM(7),CRO(96),CHL(12) , GEN(36),CLI(29), CIP(31),RIF (18)	SCC <i>mec</i> c(30)	11

	2015	<i>S.aureus</i> (50), <i>Staphylococcus</i> (168)	Cattle (200), buffalo (18)	218	ND	<i>mecA</i> (1), <i>blaZ</i> (48), <i>erm</i> (C), <i>te</i> <i>t</i> (K),(43), <i>tet</i> (M)(1)	TET(43), PEN(48),CLI, SXT	SCCm ec(1)	12
	2015	<i>E. faecium</i> (37), <i>E. faecalis</i> (13), <i>Enterococcus</i> spp.	Raw milk cheese	120(14)	ND	<i>tet</i> (M)(6), <i>tet</i> (K)(1), <i>tetL</i> (4), <i>erm</i> (B)(2), <i>aph</i> (3')(2)	CLI(14),CIP(4),KAN(11),STR(11),VAN(6),	Tn916 (7)	13
	2015	<i>S. aureus</i> (288)	Chicken(288)	288(256)	ND	<i>mecA</i> (76)	PEN(269),AMP(256),CLX(240),AMX(224),ER Y(212), TET(197),STR(150),RIF(113),AMK(99),CHL(9 1),GEN(70),CIP(39),NEL(48),SXT(39),VAN(17 )	SCCme c(76)	14
	2015	<i>S. aureus</i> (50), CONS(110)	Cattle (146), buffalo milk (14)	160(12)	ND	<i>mecA</i> (12), <i>blaZ</i> (10) , <i>tetK</i> (90), <i>tetM</i> (6), <i>er</i> <i>mC</i> (1), <i>kata</i> (6), <i>blaZ</i> (5), <i>ermB</i> (2), <i>ermC</i> ( 1), <i>tetK</i> (3), <i>tetM</i> (1)	ERY,TET, PEN	SCCm ec(12)	12
	2012	<i>S. aureus</i> (4)	dogs swab (70), cats swab (48), human nasal and oral swabs (50).	(4)	ND	<i>mecA</i> (4)	OXA(4), FOX(4), AMP(3),FOX(4),RIF(3),GEN(2),CLI(2),RIF(2),C IP(2),TET(1)	SCCm ec(4)	15
Kenya (1)	2013	<i>S. agalactiae</i> (92)	Camel(92)	92 (37)	ST617 (8), ST-612 (1),ST-616 (22)	<i>tet</i> (M) (37)	TET(37)	Tn916 (37)	16
Nigeria (5)	2018	<i>Staphylococcus</i> (24)	Beet (17), pork (3),chicken (3), Goat meat (1)	24(17)	ND	<i>mecA</i> (6), <i>tetK</i> (6), <i>mphC</i> (3), <i>er</i> <i>mt</i> (2), <i>ermC</i> (1)	FUS(19),FOX(6),OXA(6),TET(6),ERY(5),VAN( 3),SXT(2),GEN(1),STR(1),KAN(1)	SCCm ec(6)	17
	2017	<i>S. aureus</i> (30), <i>S. epidermidis</i> (16), <i>S. saprophyticus</i> (2), <i>S. sciuri</i> (1), <i>S. xylosum</i> (1)	Pork(26),beef(14) ,chicken(10)	50(48)	ND	<i>mecA</i> (49)	PEN(48),CLI(48),CHL(46),SXT(46),KAN(46),A MX(460)	SCCme c(49)	18
	2017	<i>E. faecium</i> (108), <i>E. gallinarum</i> , (30), <i>E. faecalis</i> (5), <i>E. hirae</i> . (5) <i>E. mundtii</i> (12)	Cattle (130), chickens (130),manure (130)	167 (102)	ND	<i>tet</i> (K) (NS), <i>tet</i> (L) (NS), <i>tet</i> (M) (NS), <i>tet</i> (O) (NS) and <i>erm</i> (B) (NS)	TET (102), ERY (102), CHL (13), GEN(55), STR(47),AMP(75)	ND	19

	2016	<i>E. faecium</i> (33), <i>E. casseliflavus</i> (21), <i>E. gallinarium</i> (6)	Chicken faeces (60)	(60)	ND	<i>tetM</i> (14)	ERY(60),TET(49),OFL(41),VAN(39),GEN(12)	IS256(4)	20
	2014	<i>Coagulase negative staphylococcus</i> (16)	Groin swab of dogs(16)	(16)	ND	<i>mecA</i> (16), <i>blaZ</i> (1), <i>tetK</i> (12), <i>tet(M)</i> (8), <i>erm(B)</i> (3), <i>aacA-aphD</i> (11)	PEN(16),OXA(16),FOX(16),TET(13),ERY(9),CLI(9),GEN(5),KAN(12),TOB(1),SXT(10),CHL(7)	SCCmec(16)	21
South Africa (8)	2019	<i>S.aureus</i> (120)	Poultry (120)	120(11)	ST612(11), ST36	<i>mecA</i> (12), <i>blaZ</i> (12), <i>aac(6')-aph(2'')</i> (9), <i>erm(C)</i> (3), <i>tetM</i> (10), <i>gyrA</i> (100), <i>parC</i> (580y), <i>mrs(A)</i> (3)	PEN(12),AMP(12),FOX(12),AMK(2),GEN(11),CIP(11),MXF(11),LEC(11),TET(11),ERY(9)	SCCmec(12)	22
		<i>S.aureus</i> (134)	Raw milk (134)	134	ND	<i>mecA</i> (1), <i>tetK</i> (26), <i>tetM</i> (29)	PEN(87),OXA(82),TET(48),ERY(59),AMK(24),GEN(15),CIP(7),CLI(68),CHL(4),SXT(28)	SCCmec(1)	23
	2016	<i>E. faecium</i> (180), <i>E. durans</i> (80), <i>E. hirae</i> (29), <i>E. casseliflavus</i> (20)	Cattle (340)	100	ND	<i>vanB</i> (67), <i>vanC1</i> (85), <i>vanC2/3</i> (137), <i>erm(B)</i> (137)	ERY(338),CLI(330),VAN(340),PEN(310),CET(300),STR(320),CLX(100),AMK(252),CIP(41)	ND	24
	2017	<i>S. aureus</i> (104)	Chicken(104)	(104)	ND	<i>mecA</i> (45), <i>blaZ</i> (12), <i>tet(K)</i> (32)	AMP(46),GEN(29),ERY(64),FOX(71),KAN(52),STR(57),TET(82),VAN(43)	SCCmec(45)	25
	2015	<i>S. aureus</i> (211)	Milk (211)	211 (124)	ND	<i>mecA</i> (19)	PEN (124), AMP(99), OXA (93), VAN(47), TEC(116), TET(56),ERY(56),STR(89),KAN(55),GEN(47),SXT (37)	SCCmec(19)	26
	2015	<i>E. faecalis</i> (40), <i>E. hirae</i> (100), <i>E. durans</i> (60), <i>E. faecium</i> (120)	Pigs (320)	(320)	ND	<i>vanB</i> ,(320), <i>vanC1</i> (320), <i>vanC2/3</i> (320), <i>erm(B)</i> (300)	VAN(320), STR(320) and CLX(320),STR(320),CET(286),PEN(292),CIP(248),AMO(64), AMK(272),CLI(316),ERY (280),IPM (52),	ND	27
	2014	<i>Staphylococcus spp</i> (120)	Pigs (30), Cattle (30), Cows (30), Goats (30)	120(53)	ND	<i>mecA</i> (12)	VAN(12),CRO(12),CFZ(37),CTX(19),SAM(13),PEN(53),MER(4)	SCCmec(12)	28

	2014	<i>S. xyloso</i> (18), <i>S. aureus</i> (28), <i>S. haemolyticus</i> (42), <i>S. capitis</i> (18), and other <i>Staphylococcus spp.</i> (14)	Animals (120)	(120)	ND	<i>mecA</i> (45), <i>mphC</i> (NS)	PEN (90), MER(3), VAN(14), CTX(14), CFZ(48), OXA(46), MIC(19), TET(100), ERY(14), CLI(19), NAL(120), CIP(5), OFX(6), LUX(2)	SCC <i>mec</i> <i>c</i> (45)	29
Sudan	2007	<i>S. intermedius</i> (15)	Dogs(15)	15	ND	<i>baZ</i> (17), <i>mec A</i> (14)	PEN(17),MET(14)	SCC <i>mec</i> <i>c</i> (14)	30
Senegal (1)	2012	<i>S. aureus</i> (57)	Swabs from pigs (300) and farmers	57(35)	ST5 (5),ST88(1)	<i>mecA</i> (6)	PEN(57), SXT(35), TET(20)	SCC <i>mec</i> <i>c</i> (6)	31
Tanzania (1)	2018	<i>S.aureus</i> (46), <i>CONs</i> (42)	Bovine milk(88)	88	ND	<i>mecA</i> (3)	CLI(21),VAN(2),SXT(27),TET(36),PEN(63),OX A(6),FUS(4)	SCC <i>mec</i> <i>c</i> (3)	32
	2018	<i>E.faecium</i> (77), <i>E.faecalis</i> (95), <i>E.gallinarum</i> (6), <i>E. avium</i> (6)	Cattle and cattle waste (	184(129)	ND	<i>vanA</i> (3), <i>vanB</i> (3)	AMP(4),CHL(9),GEN(117),ERY(48),RIF(129), SXT(43),TET(74), VAN(9)	ND	33
	2015	<i>E. faecium</i> (95) <i>E. faecalis</i> (9) <i>E. gallinarum</i> (7) <i>E. Hirae</i> (9)	Fecal samples of buffalo (35), wildebeest (40), zebra (40) and cattle (20)	120 (42)	ND	<i>tet</i> (W) (NS), <i>sull</i> (NS)	VAN(10),AMP(10),TET(40),SXT(32),RIF(53),E RY(42),GEN(35)	ND	34
Tunisia (10)	2018	<i>E.faecalis</i> (53), <i>E.faecium</i> (19), <i>E.casseliflavus</i> (7)	Wild birds(79)	79(37)	<i>E. faecalis</i> ST16(1),ST848(1), ST9(1)	<i>tetM</i> (33), <i>tetL</i> (12), <i>ermB</i> (22), <i>ermC</i> (5), <i>cat</i> (6), <i>aac</i> (6')- <i>le-</i> <i>aph</i> (2'')- <i>la</i> (3)	KAN(10),CHL(7),TET(37),ERY(270),CIP(170),S XT(15),PRI(3), VAN(5),STR(2),GEN(30)	ND	35
	2018	<i>S.aureus</i> (27)	Diary cows(27)	27(16)	ST97(1)	<i>mecA</i> (1), <i>blaZ</i> (13), <i>t</i> <i>etM</i> (2), <i>tetK</i> (1), <i>fusc</i> (1)	PEN(16),ERY(6),FUS(14),CLI(7),TET(7),FOX(1 )	SCC <i>mec</i> <i>c</i> (1)	36
	2017	<i>E. faecium</i> (31), <i>E. faecalis</i> (14), <i>E. durans</i> (6), <i>E. casseliflavus</i> (2), <i>E. gallinarum</i> (2)	Faecal sample of cats(20), dogs(50)	58(31)	ND	<i>erm</i> (B )(22), <i>tet</i> (M)(5), <i>tet</i> (M), <i>tet</i> (L)(16), <i>tet</i> (L)(4), <i>ant</i> (6')- <i>la</i> (11)	AMP(1),ERY(26),CIP(30), PRI(9), STR(12), KAN(12) ,GEN(9),TET(21),CHL(7)	ND	37

						, <i>aac(6')</i> - <i>le-aph(2'')</i> - <i>la</i> (16), <i>aph(3')</i> - <i>IIIa</i> (11), <i>catA</i> (1)			
2015	<i>S. aureus</i> (43)	Chicken(19), Veal(9), sheep(14), horse(1)	43(13)	ST30(1), ST398(1)	<i>tet</i> (M)(2), <i>erm</i> (C)(4), <i>erm</i> (A)(2), <i>erm</i> (T)(1), <i>tet</i> (K)(6), <i>tet</i> (L)(3), <i>tet</i> (M)(2), <i>aph(3')</i> - <i>IIIa</i> (4), <i>ant(4)-la</i> (1), <i>mrsA</i> (4)	PEN(41),OXA(2),FOX(2),KAN(4),TOB(1)	SCC <i>me</i> <i>c</i> (2)	38	
2015	<i>S. aureus</i> (17)	Goat, Cats, dogs(17)	17(7)	ST45(1),ST15(1),ST 6(1),ST2121(1),ST1 88(1)	<i>blaZ</i> (7), <i>tet</i> (M)(1), <i>e</i> <i>rm</i> (A)(1), <i>ant(6)-</i> <i>la</i> (1)	PEN(6),TET(1),ERY(1),STR(1),CIP(1)	ND	39	
2013	<i>E. faecalis</i> (49), <i>E. faecium</i> (30), <i>E. gallinarum</i> (12), <i>E. hiraе</i> (12), <i>E. casseliflavus</i> (2), <i>E. durans</i> (2)	Meat (199)	(119)	ST260(1),ST454(1), ST452(1),ST22(1),S T300(1),ST455(1),S T453(1),ST456(1)	<i>tet</i> (M) (36), <i>tet</i> (L) (32), <i>erm</i> (B) (33), <i>aac(6')</i> - <i>aph(2'')</i> (1), <i>ant</i> (6) (7)	TET(57), ERY(43), STR(17), CHLI(4),GEN (1)	ND	40	
2013	<i>E. mundtii</i> , (23) <i>E. casseliflavus</i> (20), <i>E. hiraе</i> (19), <i>E. faecalis</i> (10), <i>E. faecium</i> (10), <i>E. durans</i> (7), <i>E. gallinarum</i> (7), <i>E. dispar</i> (2)	Cattle (92)	92 (72)	ND	<i>erm</i> (B) (7), <i>tet</i> (M) (4), <i>tet</i> (L)(4)	ERY(10), TET(4) and SXT(72)	ND	41	
2012	<i>S. aureus</i> (73)	nasal swab from sheep (73)	73 (5)	ST153(5)	<i>mecA</i> (5), <i>blaZ</i> (28), <i>ant(6)-la</i> (5), <i>erm</i> (C) (5), <i>tet</i> (K) (30)	PEN(5), STR(5), KAN(5), ERY(5), TET (5), FUS(5)	SCC <i>me</i> <i>c</i> (5)	42	
2012	<i>S. aureus</i> (50)	Nasal swab of donkey(50)	50(30)	ST133(15), ST1738(4), ST1(2), ST6(4), ST2057(4), ST2110(1), ST2181(1),	<i>blaZ</i> (12), <i>erm</i> (A)(8), <i>erm</i> (C)(2), <i>tet</i> (M)(1), <i>fusC</i> (1)	PEN(12),ERY(8),TET(1),Fusic acid(12),	ND	43	

					ST1660(1)				
Uganda (1)	2017	<i>S. aureus</i> (41)	milk(30),sour milk sample(11)	41(30)	ST97(1),ST1(2)	<i>mecA</i> (23)	TET(30),RIF(1),SXT(2),ERY(1), GEN((1),CLI(1)	SCC <i>mec</i> (23)	44
Zambia	2014	<i>S.aureus</i> (31), <i>S. pseudintermedius</i> (31)	Skin(9),Ear(19), wound (2),	31(28)	ND	<i>BlaZ</i> (20), <i>tetK</i> (10), <i>tetM</i> (10), <i>ermB</i> (4), <i>tetI</i> (1)	CIP(2),ERY(5),OXA(1),PEN(28),SXT(8), TET(9)	ND	45

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