Supplementary Table 3. Geographical distribution, species, clones, and resistance mechanisms of antibiotic-resistant Gram-positive bacteria isolated from the environment in Africa from 2007-2019.

Country (n)[1]	Year	Organism/ Species (n)[2]	Specimen Sources (n)[3]	Sample size (Resistant isolates)	Clones (n)[4]	Resistance genes/ mechanisms (n)[5]	Antibiotic resistance phenotype (n)[6]	MGEs(n)[7]	Reference
Algeria	2018	*S.aureus* (33), CONS(18)	creamy cake (14), pizza (10), beef meat (10) and sausages (5)	51	ND	tetM(38), tetK(17), ermC(3), aacA-aphD(9), blaZ(25), mecA(25)	PEN(48), FOX(5), OXA(5), ERY(8), KAN(7), TET(25), VAN(1), CLI(8)	SCCmec(25)	[1]
Angola	2015	*E. faecium* (5)	Chicken farm facility (4), water from hospital and community (1)	5(4)	ST245(1), ST650(2)	*tet*(M)(4), *erm*(B)(4), *tet*L(2)	TET(4), ERY(4), STR(4), NIT(2), Q/D	ND	[2] 4
Egypt	2016	*S. aureus* (23)	Food sample (23)	23(NS)	ST689(1)	*mecA*(3), *van* A(1), *van*B(1)	VAN(NS), CIP(NS), GEN(NS), SXT(NS), OXA(NS), ERY(NS)	SCC*mec*(3)	[3]
Ghana	2018	*S.aureus* (5)	Ward (5)	(5)	ST15(4), ST508(1)	*BlaZ(5), dfrG(2)*	PEN(5)	ND	[4]
Nigeria (1)	2017	*E. faecium (100)*	Vegetables soil, farm, Cloacal swabs (25), Manure (8), Rectal swabs (2)	(100)	ND	*aac(6')-Ie-aph(2")-Ia*(35), *aph(2')-1c*(31) ,*aph(3')-IIIa*(32), *ant(4')-Ia(14)*	AMP (63), GEN(37)	ND	[5]

[1] Total number of studies per country
[2] Total number of isolates
[3] Total number of specimen source
[4] Total number of resistant clones
[5] Total number of resistant clones
[6] Number of different antibiotics any one isolate is resistant to
[7] Total number of mobile genetic elements : plasmids,transposons, integrons

South Africa (4)	2019	*E.faecalis (65), E.faecium(45)*	Sewage water (56), treated effluent (19), hospital waste (32)	110	*E. faecium* ST25, ST23, E.faecalis ST23, ST25,ST780	*ermB(48), tetM(47), tetL(44), vanC(9),aph(3”) -IIIa(47), aac(6')-Ie-aph(2”)-Ia(27),*	ERY(48), TET(47), VAN(9)	ND	6
	2019	*E.faecium(38), E.faecalis(17)*	Ground and surface water (56)	56	ND	*VanA(9), vanB(2)*	VAN(44), STR(17), CHL(17), AMP(25), TET(28), GEN(9) SMX(9)	ND	7
	2017	*S. aureus* (30)	Recreational waters and beach sand (30)	(30)	ND	*mecA*(5), *fem*A(16). *rpoB*(11), *blaZ*(16),e rm*B*(15), *tet*(M)(8)	AMP (29), PEN (29), RIF(24), CLI(24), OXA (22), ERY(21), VAN(15), TET(13), SXT(13), CIP(10), GEN(1)	SCCmec(5)	8
	2016	*E. faecium* (30), *E. faecalis* (37) *E. mundtii* (36), *E. casseliflavus* (14), *E. gallinarum* (5), *E. hirae*(1), *E. sulfurens*(1)	Surface water(124)	124(86)	ND	*tet*(L)(17), *msrC*(9)	AMP(59), AMX(53), PEN(87), STR(8), VAN(86), CHL(23), CIP(47), ERY(68), TET(59)	ND	9
	2015	*E. faecium* (30), *E. durans.* (15)	waste water (32) and effluent (32)	(45)	ND	*erm*(B) (40), *vanB,* (42), *vanC1* (42), *van C2/3* (42)	PEN (38), ERY(40), CTX(43), GEN(28), IPM(43), TET(45), KAN(43), CIP(43), VAN(29), CLI(45)	ND	10

	2017	*S. aureus* (12)	Wastewater	12	ST3245 (7), ST15 (1)	*bla*Z(7), *msr*A(7), *tet*(K)(1)	PEN(12), ERY(7), TET(1), CLI(1)	ND	[12]
	2017	*E. faecalis* (2), *E. faecium* (NS), *Enterococcus spp* (NS)	Urban wastewater (5)	5(2)	ST86(2)	*optr*A(2), *erm* (A), *erm*(B), *tet*(M)(1), *tet*(L)(1), aac(6')-aph(2"),	CHL(2), CIP(2), ERY(2), TET(1), GEN(1), STR(2)	ND	[13]
Tunisia (7)	2016	*E. faecium (86), E. faecalis*(8), *E. casseliflavus* (6)	Hands (50), inanimate such as beds, treatment tables, toilets, faucets, wrists, sinks (250)	(100)	ST910 (13), ST80 (1)	*erm*(B) *(71), tet*(M) (18), *aph(3')-IIIa* (27), *ant(6)-Ia (15),cat*(A) (4), *vanC2*(6)	ERY(73), TET(20), STR(27) and KAN(28), VAN(14), CHL(10), SXT(100), CIP(48), PRI(18)	*IS16* (14)	[14]
	2016	*S. saprophyticus* (30), *S. haemolyticus* (38), *S. epidermidis* (NS), *S. cohnii* (NS), *S. warneri* (NS), *S. sciuri* (NS), *S. simulans* (NS), *S. pasteuri* (NS), *S. arlettae* (NS) and *S. xylosus* (NS)	Inanimate surfaces (83	83 (32)	ND	*mecA*(20), *msr*(A)(32), *erm*(C)(8), *tet*(K)and/or *tet*(M)(21), *aac*(6´)-Ie-*aph*(2´´)-Ia (16),(*aph*(3`)-IIIa(19), *ant*(4´)-Ia (14), *ant*(6´)-Ia (3)	ERY(32), TET(23), GEN(16), TOB(14), STR(3), Pen(39), OXA(20), FOX(20), KAN(27), SXT(17), CLI(14), CIP(18), FUS(26), STR(11)	SCC*mec*(20)	[15]
	2015	*E. faecium* (34), *E. hirae* (23), *E. faecalis (4)*, and *E. casseliflavus* (4)	Vegetable food (34), soil and irrigation water (27)	65 (40)	*ST2 (5), ST16 (2), ST528 (2), ST56 (1), ST885 (1), ST886 (1)*	*erm*(B) (12), *tet(M)-tet(L)(10), aph(3')-III, (10) ant(6) (2),vanC2(4)*	CIP(42), ERY(12), TET(10), KAN(10), CHL(5), STR(2), and GEN(5), VAN(4)	ND	[16]

	2015	*E. faecium* (54), *E. faecalis* (17), *E. hirae* (8), *E. casseliflavus* (4), *E. durans* (2)	waste and surface water (114)	(85)	ST480 (1), ST531 (1), ST55 (1), ST532(1), ST202 (1), ST314(1), ST985(1), ST30 (1), ST986 (1), ST12 (1), ST296 (1), ST327(1)	*aph(3')-IIIa* (22), *ant(6)-Ia (4), erm*(B) (34), *tet*(M) (13), *tet(L)(8),aac(6')-Ie-aph*(2')(15)	GEN(22), KAN(22), STR(7), ERY(36), TET(13), SXT(79), CIP(6),	ND	[17]
	2015	*S. aureus* (12)	Hospital environment (12)	12(6)	ST247(2)	*bla*Z(12), *erm*(A), *tet*(M)(2), *aac(6')-aph(2')*(2), *mec*A(2)	STR(2), KAN(2), ERY(2), CLI(2), TET(2), FUS(2), TOB(2), GEN(2), AMK(2), OXA(6), PEN(12), FOX(2)	SCCmec(2)	[18]
	2014	*E.faecium* (5), *E. casseliflavus* (7)	Hospital environment ((beds, treatment table, toilet, faucet, wrist and sink) (100)	(12)	ST80 (1)	*van*A(5), *van*C2(7), *erm*B(12), *tet*M(5), *aph(3')-IIa*(5), *aac(6')-aph(2'')*(5)	VAN (12), AMP(5), CIP(12), ERY(12), TET(8), STR(6), KAN(80, SXT(11), GEN(3), TEC(5)	*IS16(1)*	[19]
	2014	*S. aureus (10)*	Refrigerator, floor, desk, sink, tray surface (10)	10(5)	ND	*BlaZ(5), tetK(5), tetM(2),*	SXT(5)	*ND*	[20]
Zambia									

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