

グラム陽性球菌

	PCG	ABPC	CEZ	CDTR-PI	CZOP	IPM/CS	CVA/AMPC	GM	ABK	CAM	CLDM	MINO	VCM	FOM	LVFX	ST
MRSA	0.0	0.0	0.0	0.0	0.0	0.0	0.0	71.4	97.1	14.5	70.0	92.9	100.0	84.7	10.5	99.6
MSSA	43.5	43.5	100.0	99.8	100.0	99.9	100.0	77.9	99.1	74.2	93.5	99.7	100.0	95.5	82.6	99.9
Staphylococcus epidermidis	23.2	23.2	54.5	54.5	54.5	54.6	57.8	56.2	99.6	59.0	74.6	96.1	100.0	55.4	44.6	97.7
Enterococcus faecalis	99.7	99.7	0.0	0.0	0.0	100.0	100.0	0.0	0.0	6.5	0.2	30.4	100.0	0.2	89.9	0.0
Enterococcus faecium	0.0	0.0	0.0	0.0	0.0	0.8	0.0	0.0	0.0	0.0	6.2	29.4	100.0	0.8	12.5	0.0

腸内細菌

	ABPC	PIPC	CEZ	CTM	GTRX	CFPM	CAZ	GMZ	IPM/CS	MEPM	AZT	SBT/ABPC	TAZ/PIPC	GM	AMK	MINO	FOM	LVFX	ST
Escherichia coli	70.2	74.3	84.3	88.0	88.2	88.8	88.4	99.9	100.0	100.0	88.3	81.1	99.7	94.9	100.0	96.3	91.7	75.0	90.1
Klebsiella pneumoniae	0.2	56.9	94.6	96.1	96.3	97.2	97.2	100.0	99.5	99.5	96.6	82.8	98.3	98.0	100.0	93.2	2.9	97.6	96.1
P.mirabilis	83.5	87.8	87.4	87.8	89.0	90.9	92.5	99.2	99.2	100.0	88.2	88.2	100.0	74.5	99.6	40.0	56.4	68.2	96.4
Serratia marcescens	29.2	77.7	0.0	14.6	78.4	99.6	86.8	89.4	99.6	100.0	86.1	28.1	87.9	99.2	99.6	94.8	64.6	96.3	99.2
M.morganii	3.1	73.4	1.0	13.8	75.5	88.3	91.4	95.7	15.9	100.0	81.9	17.0	100.0	97.8	100.0	54.2	2.1	82.9	98.9
E.aerogenes	19.4	75.9	8.3	43.5	72.2	99.0	81.4	8.3	70.3	100.0	87.9	62.9	93.5	100.0	100.0	95.3	46.3	96.3	100.0

ブドウ糖非発酵菌

	PIPC	CFPM	CAZ	IPM/CS	MEPM	AZT	TAZ/PIPC	GM	AMK	MINO	FOM	GPFX	LVFX	ST
Pseudomonas aeruginosa	96.8	97.3	97.7	80.2	84.8	89.9	96.6	78.4	97.9	0.0	0.4	92.3	90.1	0.0
Acinetobacter	45.1	93.2	97.6	100.0	100.0	28.3	98.5	86.5	96.6	55.7	0.9	31.2	43.2	35.1
Stenotrophomonas maltophilia	47.3	13.5	44.5	0.0	0.0	4.0	52.7	59.4	68.9	98.6	1.3	25.6	72.9	100.0

呼吸器関連菌

	PCG	ABPC	CTX	CTRX	CDTR-PI	CFPN-PI	FRPM	CAM	CLDM	MINO	LVFX
Streptococcus pneumoniae	50.8	50.8	97.1	95.9	99.0	98.0	98.3	29.0	48.2	51.0	98.3

	ABPC	OCL	CTX	CTRX	CDTR-PI	MEPM	FRPM	CVA/AMPC	CAM	MINO	CP	LVFX
Haemophilus influenzae	27.8	42.0	100.0	100.0	100.0	98.1	58.7	66.7	83.1	98.4	99.9	99.9

	PCG	ABPC	CTX	CTRX	CDTR-PI	CFPN-PI	CAM	CLDM	MINO	VCM	LVFX
Streptococcus group A	100.0	100.0	100.0	100.0	100.0	100.0	60.4	74.4	88.3	100.0	90.7