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# Evaluation of the most common microorganisms causing surgical site infections after renal transplantation

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# Background

Renal transplant recipients are highly susceptible to infections due to surgical complexity and immunosuppression, with surgical site infections (SSI) being a significant early postoperative complication.

This study aimed to identify the common microorganisms causing SSI after renal transplantation and evaluate their susceptibility to standard antibiotics.

# Methods

- Single-center, observational cohort study of adult renal transplant patients from January 2017 to December 2019 (n=231)
- Cephalosporins were used for prophylaxis
- Evaluation of risk factors for SSI (age, gender, body mass index (BMI), type of donation) and clinical outcomes (organ function, patient survival, SSI occurrence, hospital stay)
- Patients were grouped by BMI (<25, 25-30, >30)
- Microbial specimens from SSI cases (n=46) were analyzed using conventional methods and antibiotic sensitivity tests

# Results

231 kidney transplant recipients ( $56 \pm 12.8$  years, male 149, living donation 41)

46 developed SSIs ( $55 \pm 12,82$  years, male 31, living donation 11)

SSIs occurred in 15.4% of BMI group 1, 18.5% of group 2, and 28.8% of group 3

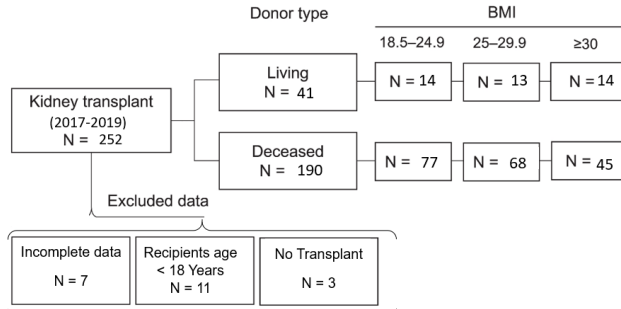
Of the 46 SSIs, 29 (63%) were monomicrobial, and 17 (37%) were polymicrobial

Predominant organisms: Staphylococcus epidermidis (23.9%), Enterococcus faecalis (21.7%), coagulase-negative staphylococci (26.1%), and Candida albicans (21.7%)

Numerous other bacteria and fungi against which the routine antibiotic was not effective, were found, especially in patients with a higher BMI

# Results

## Patient flow



## Conclusion

SSI is an early complication after renal transplantation, which brings a great burden to patients. Therefore, preventing wound infection is extremely important. Routinely used anti-infective substances do not cover all bacteria and fungi, which especially occur in patients with a higher BMI. Therefore, **the routine anti-infective therapy should be re-evaluated in further studies and possibly expanded in high-risk patients.**

## Organisms causing SSI

	Organismus	Relative Häufigkeit (%) (n = 14 Organismen)
<b>Gramm positiv</b>	Staphylokokkus epidermidis	11 (23,91%)
	Staphylokokkus haemolyticus	3 (6,52%)
	Staphylokokkus Warneri	1 (2,17%)
	Enterokokkus faecium	6 (13,04%)
	Enterokokkus faecialis	10 (21,74%)
	Lactobacillus spp.	3 (6,52%)
	<b>Gramm negativ</b>	Koagulase-negative Staphylokokken
Klebsiella spp.		4 (8,70%)
Acinetobacter lwoffii		2 (4,35%)
Bacteroides thetaiotaomicron		2 (4,35%)
Enterobacter cloacae complex		1 (2,17%)
Enterobacter Hormaechei		1 (2,17%)
<b>Fungi</b>	Candida albicans	10 (21,74%)
	Candida spp.	4 (8,70%)

