

A blind spot in an FDA standard leaves significant bacterial survivors after chlorhexidine gluconate preoperative skin preparation

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BACKGROUND:

- Skin-flora bacteria are the most significant contributor to surgical site infections (SSIs)¹
- Presurgical skin preparation (PSP) is the cleansing of skin prior to surgery, usually with alternating scrubs of antiseptic and alcohol
- The FDA requires PSP products to demonstrate a 2-3 log₁₀ reduction of natural flora using a non-destructive skin sampling technique known as the Cup Scrub method²
- We developed a porcine model for PSP testing using a microbiological sampling technique known as the Tissue Blend method³
- We hypothesized that on-label use of antiseptics approved by the Cup Scrub method would leave viable bacteria underneath the skin's surface. To test our hypothesis, we quantified and characterized the bacterial survivors after applying chlorhexidine gluconate (CHG).

METHODS: We applied the Cup Scrub and Tissue Blend methods to the backs of 7 Yorkshire pigs (Figure 1) following alternating scrubs of 4% CHG and alcohol (n=5 sites/pig). Control skin was used as a baseline (n=5 sites/pig). With 4 treatment groups, 20 samples were taken from each animal for a total of 140 samples.

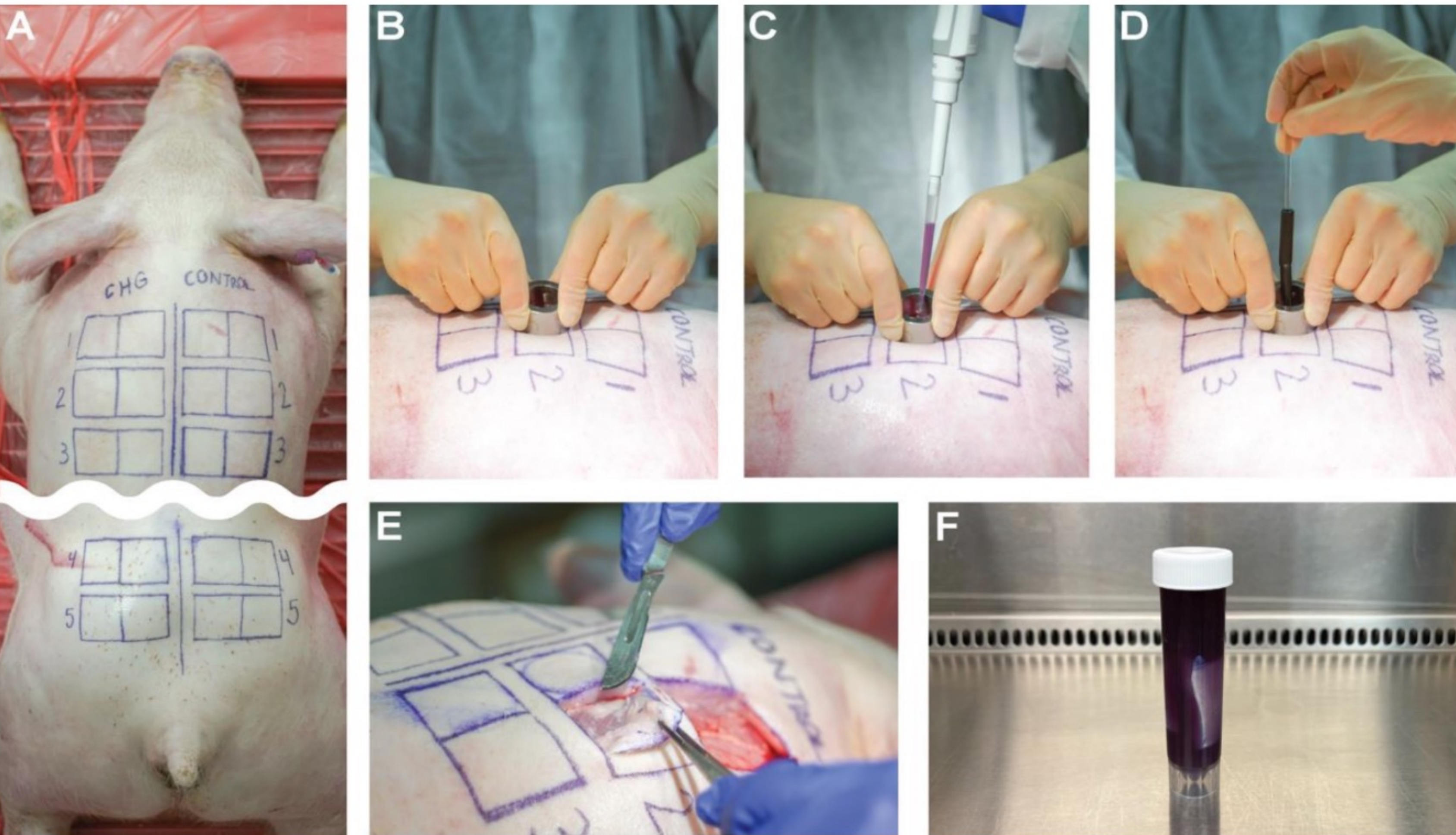


Figure 1: Location and process of sample collection. **(A)** Pig back marked with sample location boundaries. **(B-D) Cup Scrub Method:** A sterile containment cylinder is placed on the skin after applying a PSP. Neutralizing broth is pipetted into the cylinder; the skin is then agitated with a rubber spatula to suspend bacteria into solution. **(E-F) Tissue Blend method:** Full-thickness skin samples were excised, homogenized, serially diluted, and plated on agar.

- RESULTS:**
- Cup Scrub Method log₁₀ reduction = 1.57 +/- 0.45 CFU/cm² (Initial = 2.62 +/- 0.21 log₁₀ CFU/cm²)
 - Tissue Blend Method log₁₀ reduction = 0.23 +/- 0.48 CFU/cm² (Initial = 3.46 +/- 0.24 log₁₀ CFU/cm²)
 - Bioburden grouped by location zone was similar across anatomical sites along the pig back.
 - Vast majority of bioburden were Gram-positive cocci. Most were from the *Staphylococcus* family

