

Supporting information:

Time-gated luminescent in-situ hybridization (LISH): highly sensitive detection of pathogenic *Staphylococcus aureus*

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HPLC data of conjugated oligonucleotides:

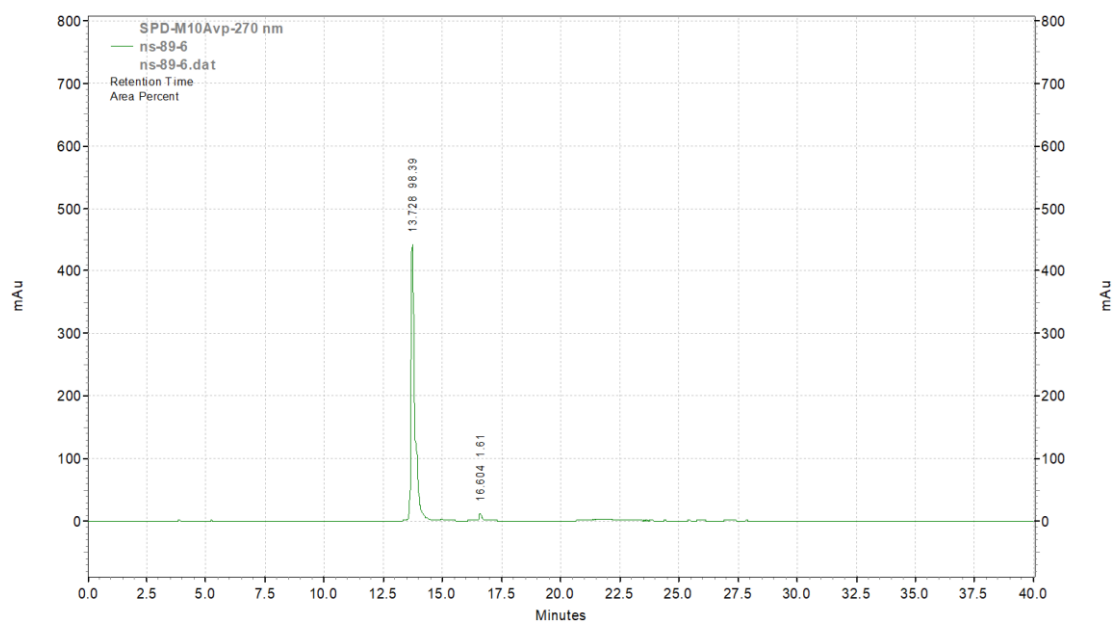


Figure S1: HPLC chromatogram of SAU69 DNA at 270 nm (Purity 98.4%, RT=13.7).

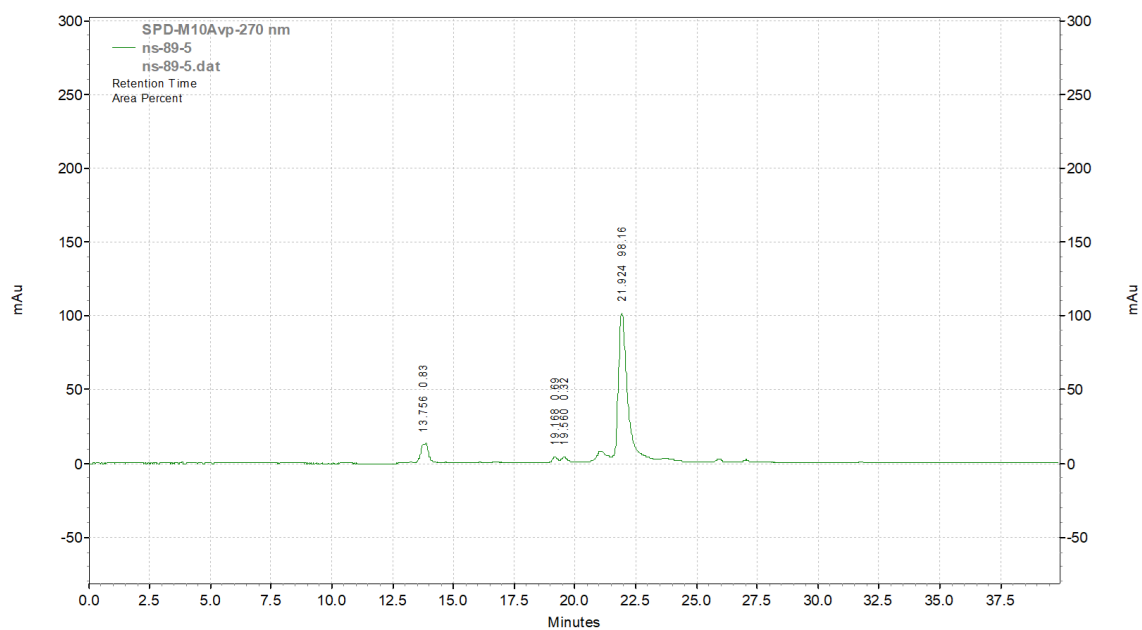


Figure S2: HPLC chromatogram of SAU69-BHHTEGT probe at 270 nm (Purity 98.1%, RT=21.9).

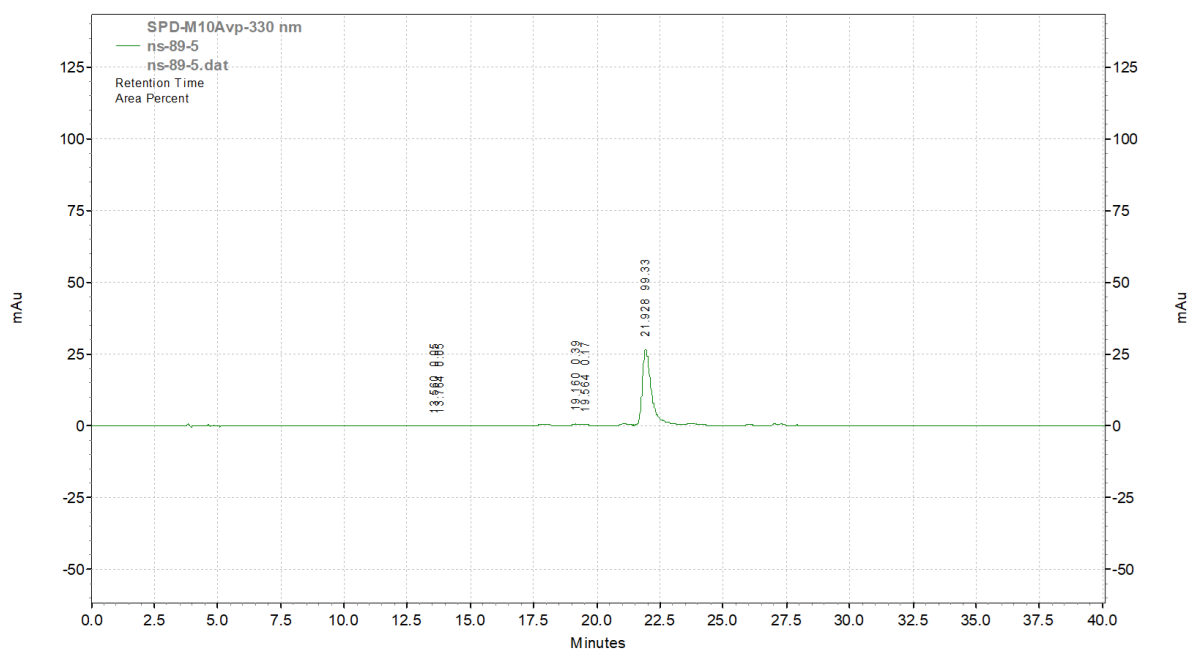


Figure S3: HPLC chromatogram of SAU69-BHHTEGST Probe at 330 nm (Purity 99.3%, RT=21.9).

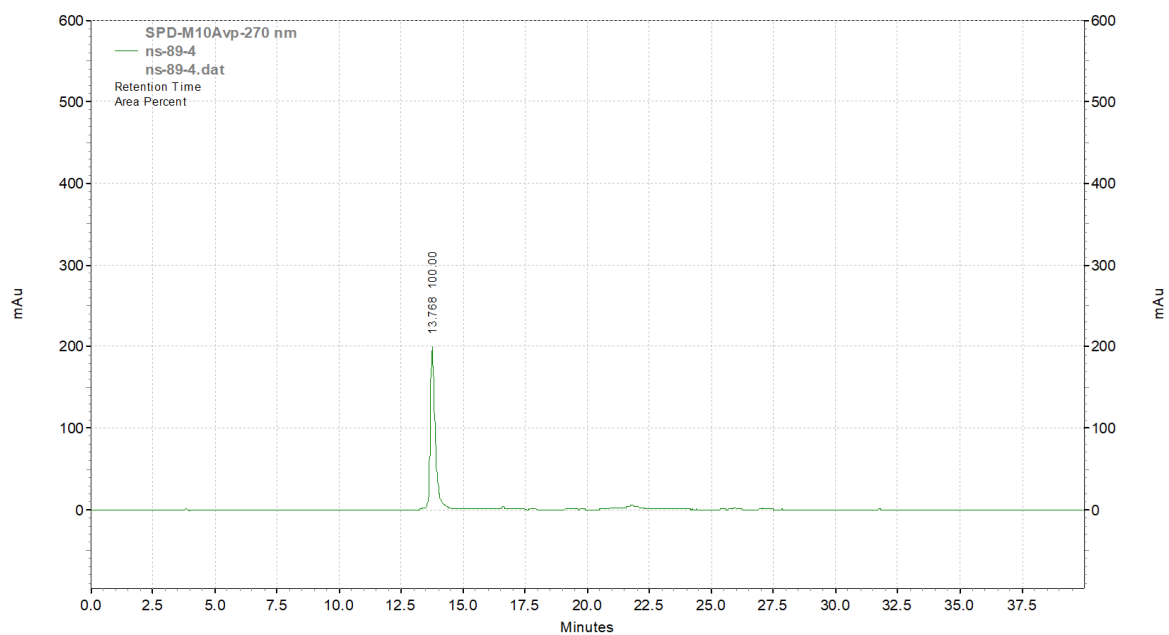


Figure S4: HPLC chromatogram of EUB338 DNA at 270 nm (Purity 100%, RT=13.77).

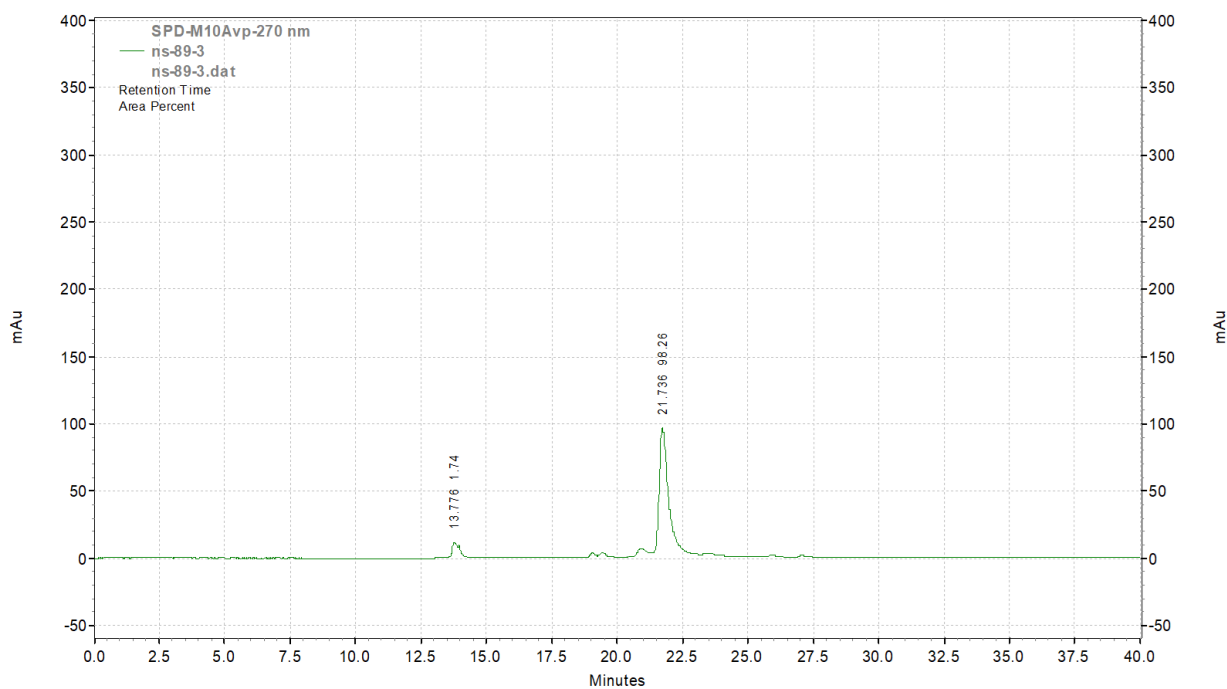


Figure S5: HPLC chromatogram of EUB338-BHHTEGST Probe at 270 nm (Purity 98.2%, RT=21.7).

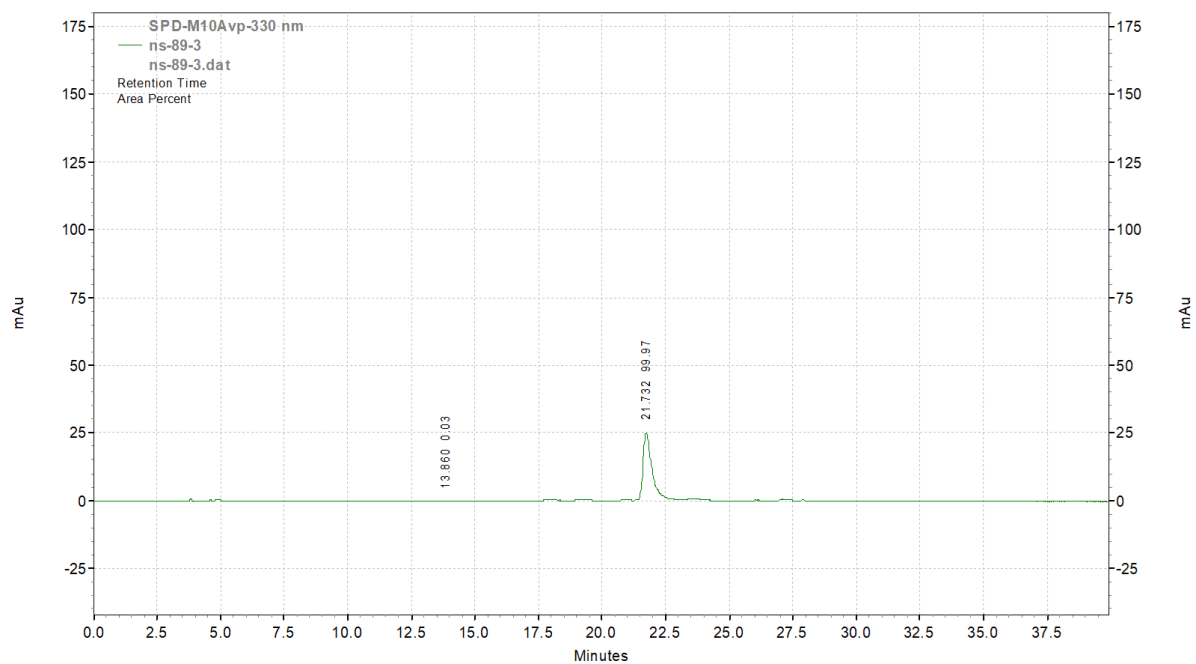


Figure S6: HPLC chromatogram of EUB338-BHHTEGST Probe at 330 nm (Purity 99.9%, RT=21.7).

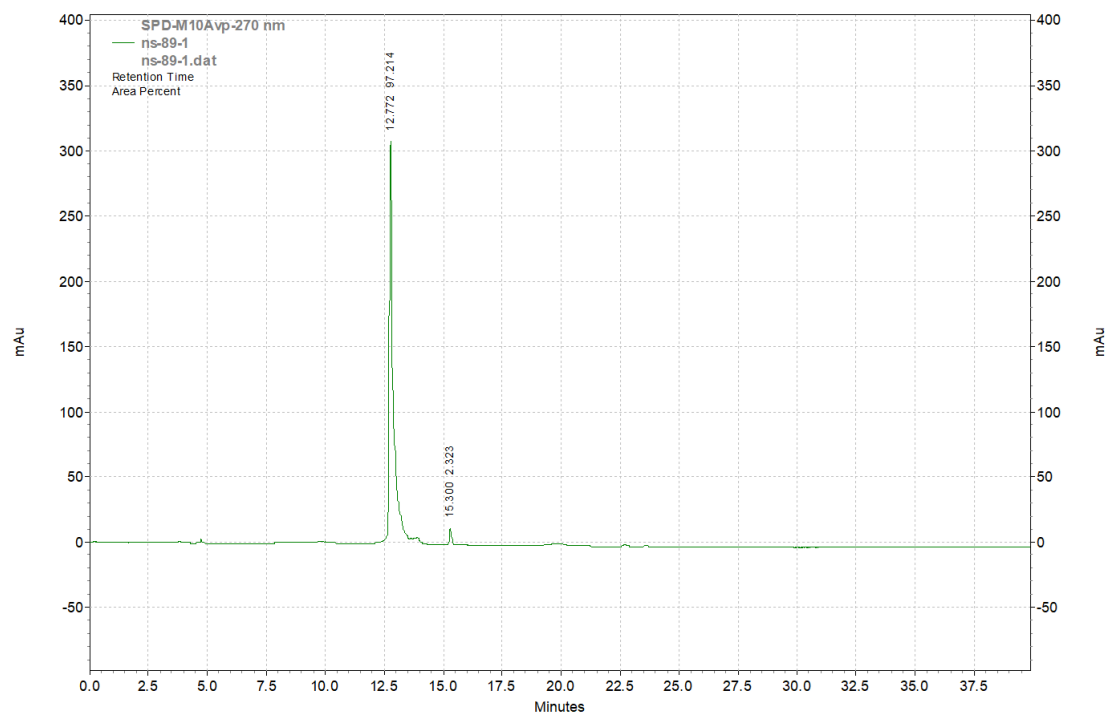


Figure S7: HPLC chromatogram of NON-EUB338 DNA at 270 nm (Purity 97.2%, RT=12.77).

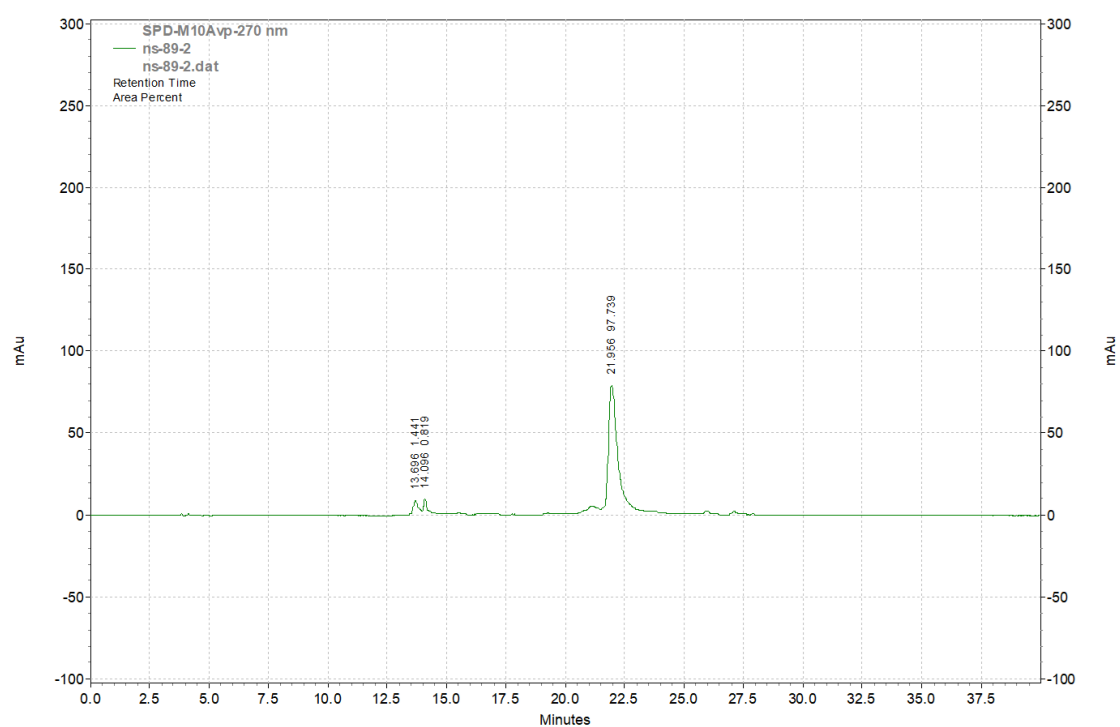


Figure S8: HPLC chromatogram of NON-EUB338-BHHTEGT Probe at 270 nm (Purity 97.7%, RT=21.96).

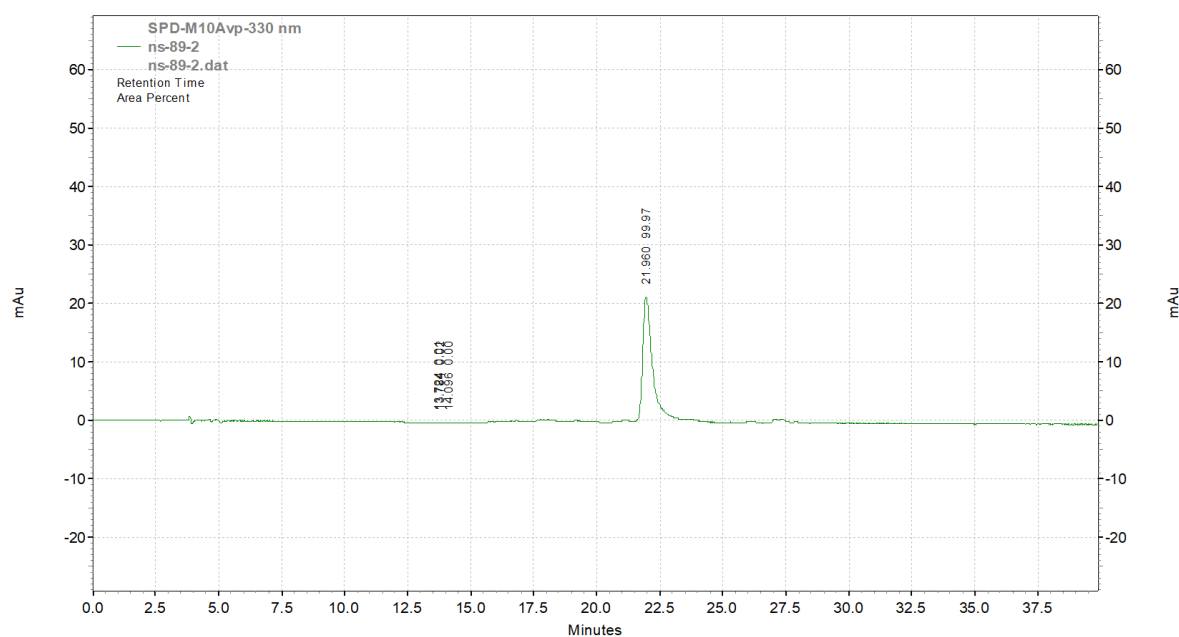


Figure S9: HPLC chromatogram of NON-EUB338-BHHTEGST Probe at 330 nm (Purity 99.97%, RT=21.96).

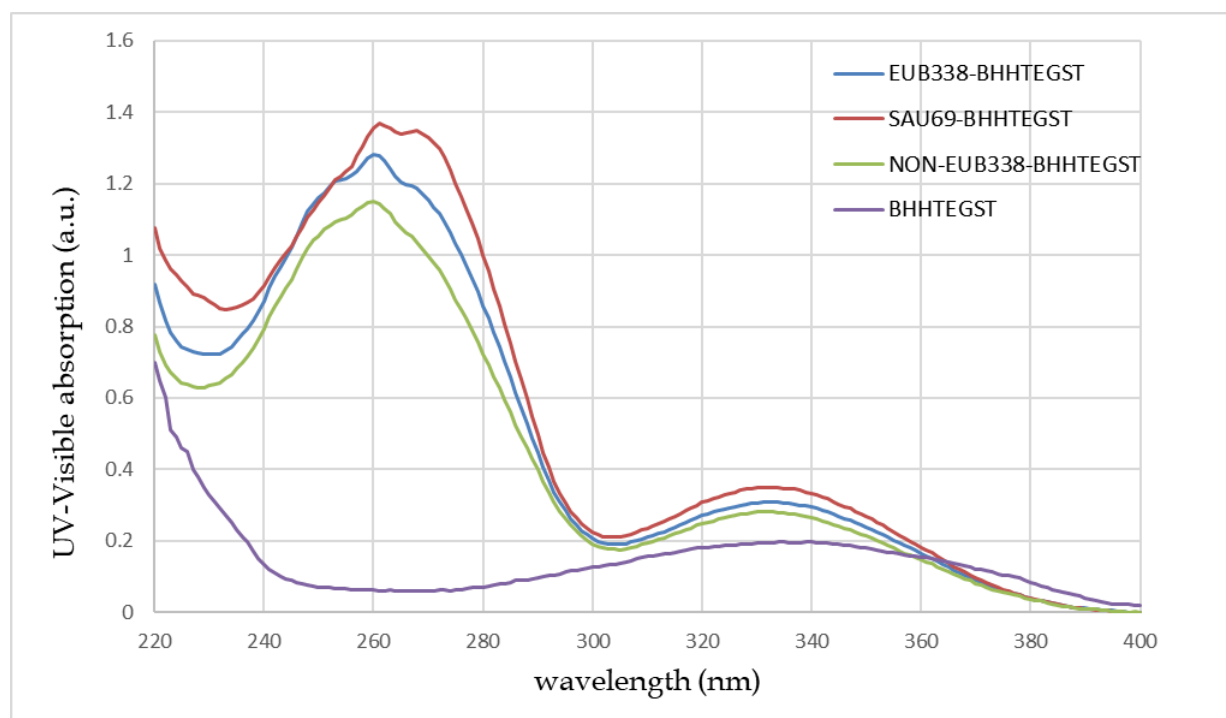


Figure S10: UV-visible absorption spectra for the BHHTEGST ligand and conjugated oligonucleotide probes ($\sim 70 \mu\text{M}$) (BHHTEGST $\lambda_{\text{ex,max}}$ = 335 nm and oligo at 260 nm).

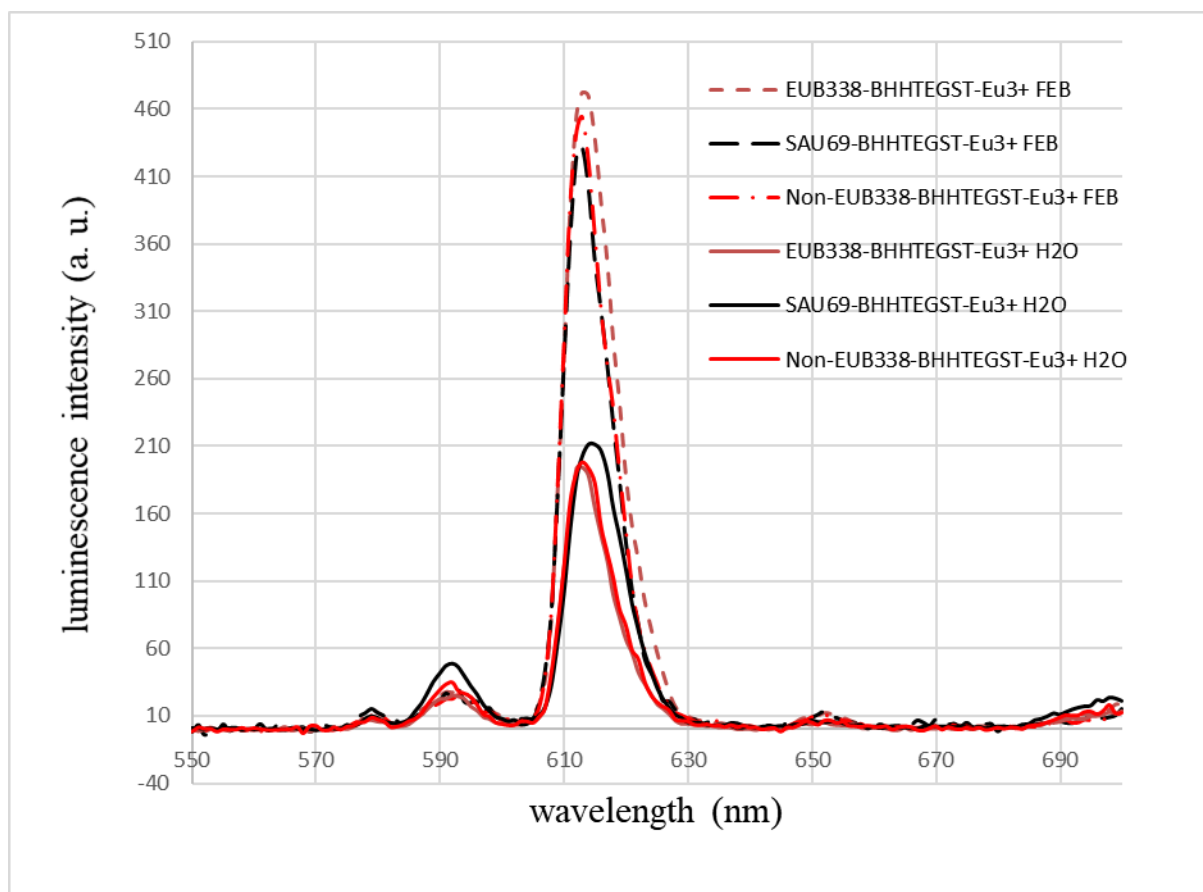
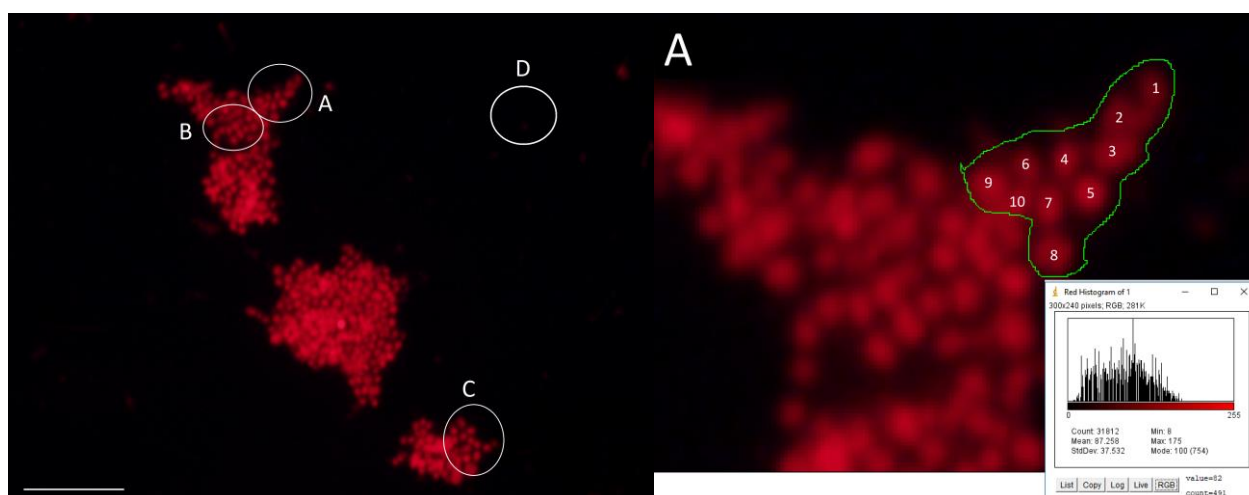


Figure S11: Luminescence emission of oligo conjugates (DNA-BHHTEGST-Eu³⁺ DNA=EUB338, SAU69 and Non-EUB338 (5.0 μ M) in FEB and MQ water. Excited at 335 nm and maximum luminescence emission observed at 613 nm.



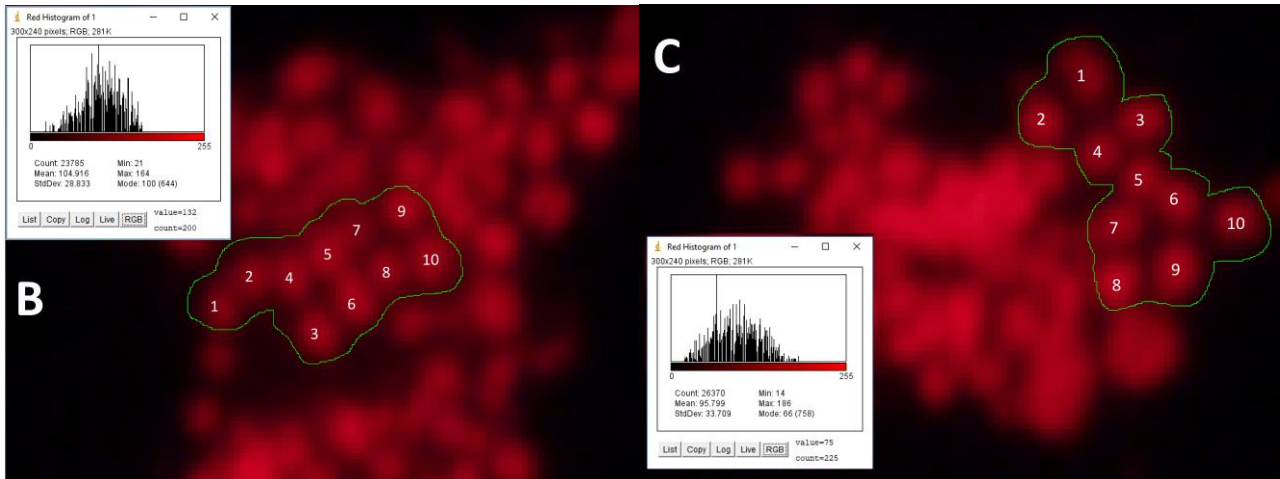


Figure S12. Quantification of the signal-to-noise ratio for the SAU69-BHHTEGST-Eu³⁺ probe LISH labelled *S. aureus* using time-gating is shown. The circles (A, B and C) show the regions used for the quantification of the mean signal intensity of the desired target cells and (D) for the background noise; (A, B & C) [zoomed in images] Average mean signal 95.7 [Mean brightness of 10 cells in histogram red channel]; (D) the Background 3.1 [mean brightness of region of background in histogram red channel (not shown)] were measured. The signal-to-noise ratio (SNR) was calculated to be 31.

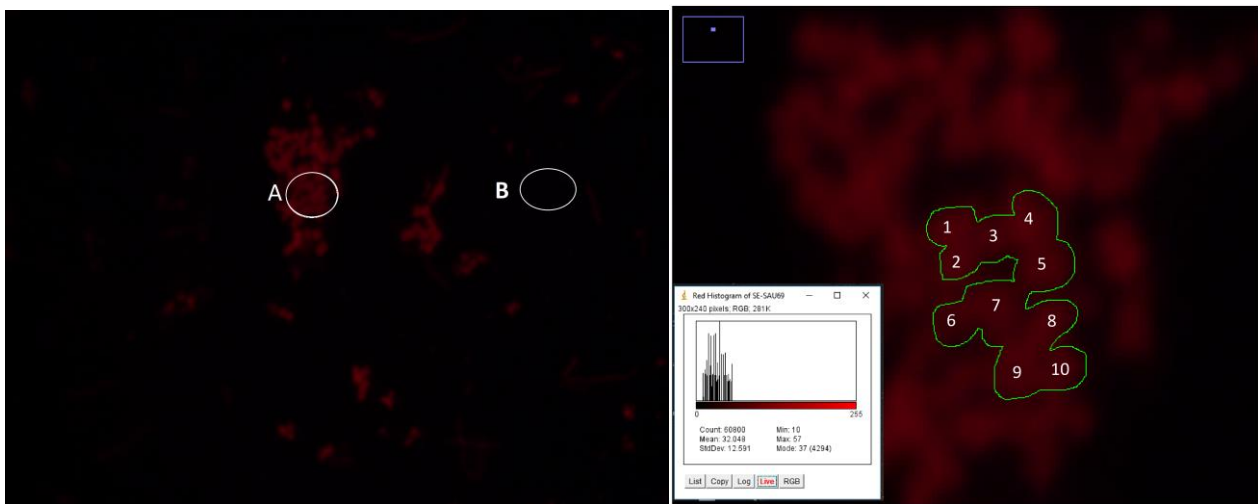


Figure S13. Quantification of the signal-to-noise ratio for the SAU69-BHHTEGST-Eu³⁺ probe LISH labelled *S. epidermidis* using time-gating conditions is shown. The circles (A) show the regions used for the quantification of the mean signal intensity of the desired target cells and (B) for the background noise; (A) [zoomed in images] Mean signal 32 [Mean brightness of 10 cells in histogram red channel]; (B) the Background 6.9 [mean brightness of region of background in histogram red channel (not shown)] were measured. The signal-to-noise ratio (SNR) was calculated to be 4.6.

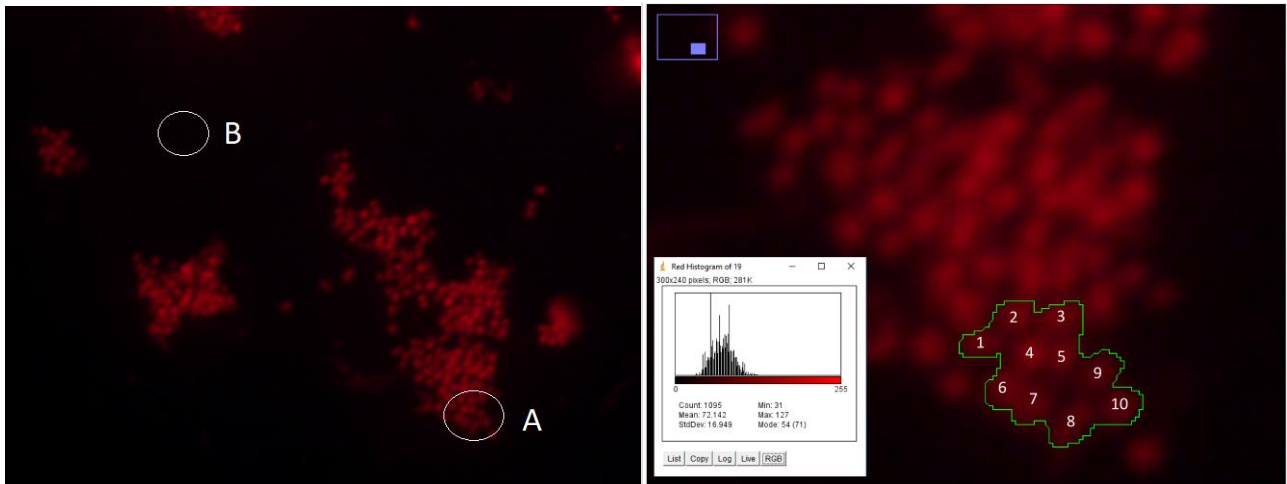


Figure S14. The quantification of the signal-to-noise ratio in EUB338-BHHTEGST-Eu³⁺ probe LISH (Top image) labelled *S. aureus* collected under time-gated condition is shown. The circles (A) show the regions used for the quantification of the mean signal intensity of the desired target cells (n=10) and (B) for the background noise; (A) [zoomed in images] Mean signal 72.1 [Mean brightness of 10 cells in histogram red channel]; (B) the Background 5 [mean brightness of region of background in histogram red channel (not shown)] were measured. The signal-to-noise ratio (SNR) was calculated to be 14.4.

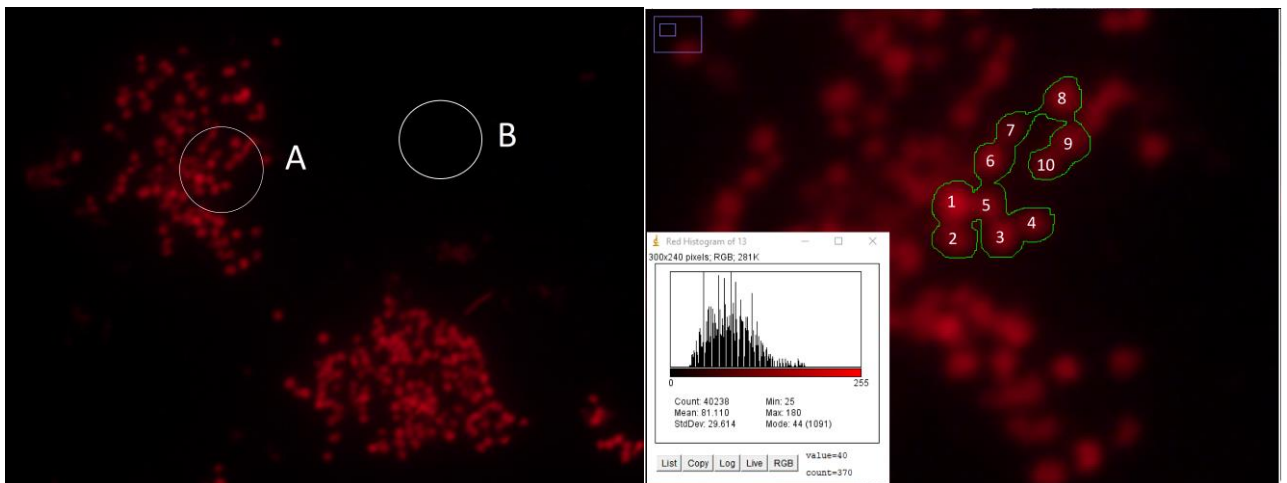


Figure S15. The quantification of the signal-to-noise ratio in EUB338-BHHTEGST-Eu³⁺ probe LISH (Top image) labelled *S. epidermidis* collected under time-gated condition is shown. The circles (A) show the regions used for the quantification of the mean signal intensity of the desired target cells (n=10) and (B) for the background noise; (A) [zoomed in images] Mean signal 81 [Mean brightness of 10 cells in histogram red channel]; (B) the Background 5 [mean brightness of region of background in histogram red channel (not shown)] were measured. The signal-to-noise ratio (SNR) was calculated to be 16.2.

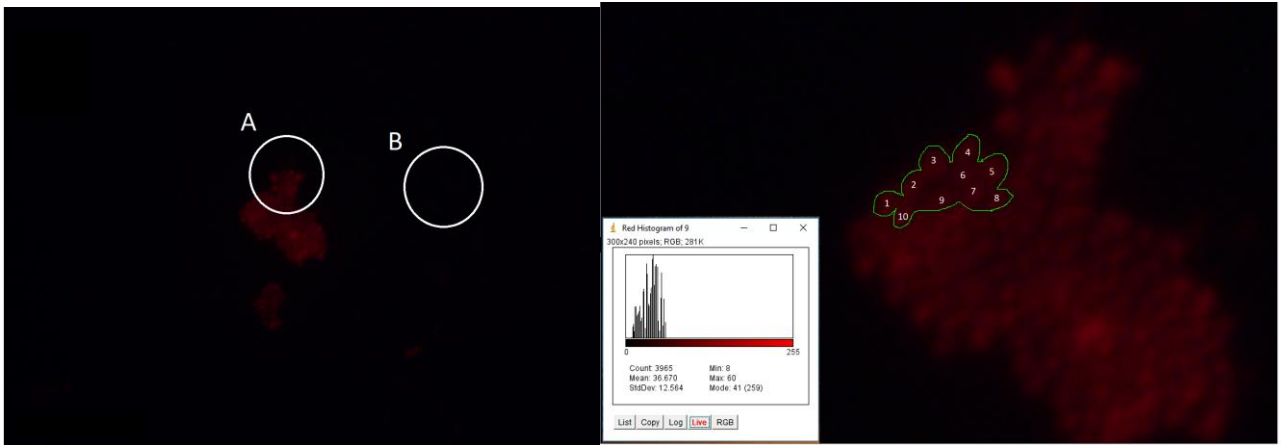


Figure S16. The quantification of the signal-to-noise ratio in Non-EUB338-BHHTEGST-Eu³⁺ probe LISH (Top image) labelled *S. aureus* collected under time-gated condition is shown. The circles (A) show the regions used for the quantification of the mean signal intensity of the desired target cells (n=10) and (B) for the background noise; (A) [zoomed in images] Mean signal 36.7 [Mean brightness of 10 cells in histogram red channel]; (B) the Background 6.1 [mean brightness of region of background in histogram red channel (not shown)] were measured. The signal-to-noise ratio (SNR) was calculated to be 6.

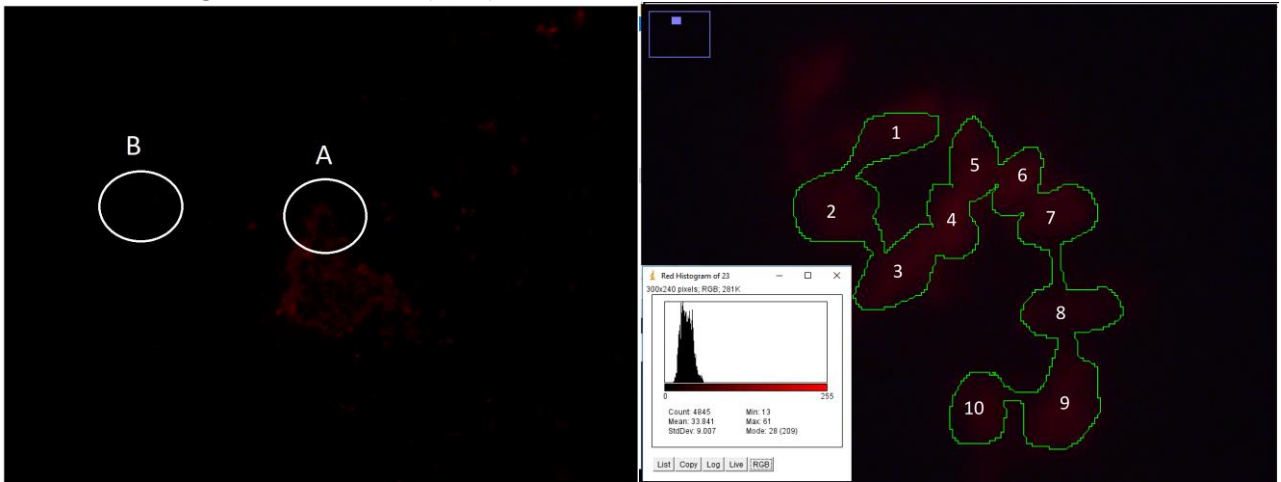


Figure S17. The quantification of the signal-to-noise ratio in EUB338-BHHTEGST-Eu³⁺ probe LISH (Top image) labelled *S. epidermidis* collected under time-gated condition is shown. The circles (A) show the regions used for the quantification of the mean signal intensity of the desired target cells (n=10) and (B) for the background noise; (A) [zoomed in images] Mean signal 33.8 [Mean brightness of 10 cells in histogram red channel]; (B) the Background 7.3 [mean brightness of region of background in histogram red channel (not shown)] were measured. The signal-to-noise ratio (SNR) was calculated to be 4.6.

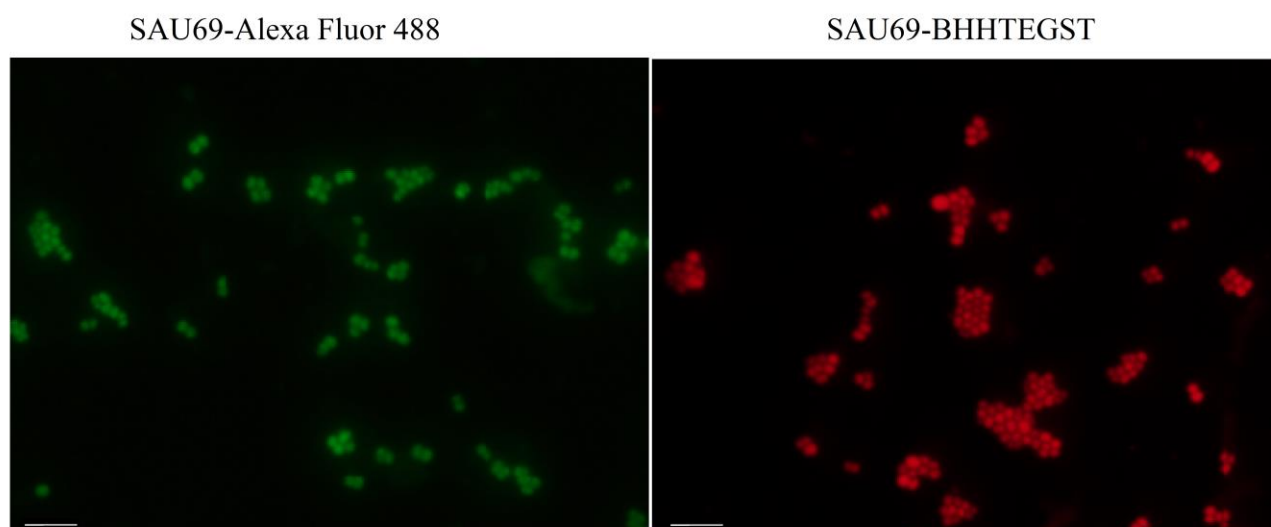


Figure S18. Staining of *S. aureus* cells with SAU69-Alexa Fluor 488 probe (left) and time gated luminescence (TGL) imaging SAU69-BHHTEGST staining (right). Scale bar 5 μ m.

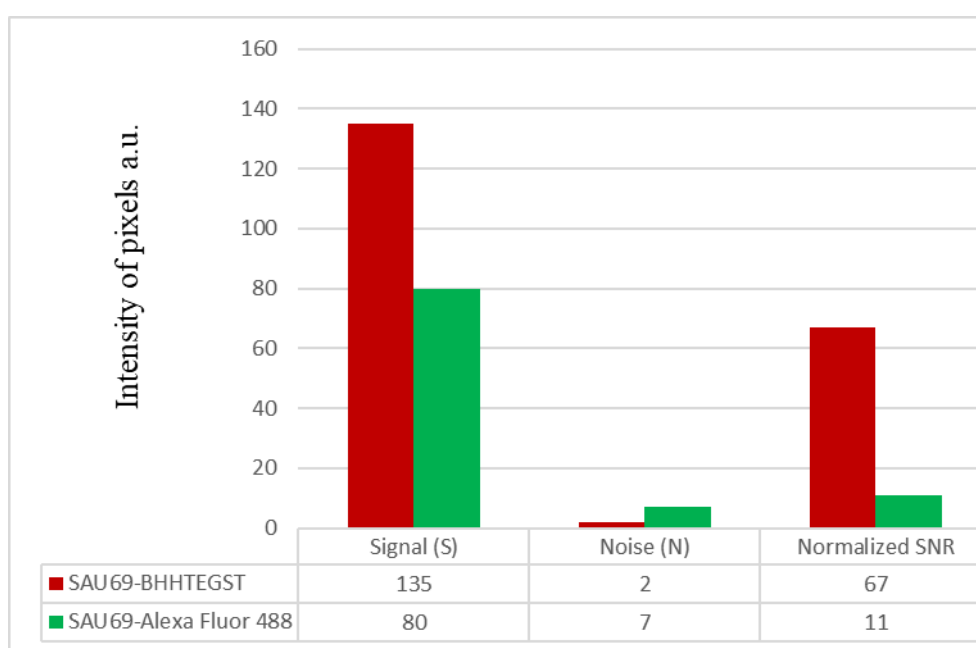


Figure S19. Representation of signal-to-noise ratios (SNRs) for *S. aureus* cancer cells labeled with SAU69-BHHTEGST and SAU69-Alexa Fluor 488.

Table S1: Results of images in Figure S10 to S15; The mean signal and background staining of *S. aureus* and *S. epidermis* are shown. These were labelled with SAU69-BHHTEGST-Eu³⁺, EUB338-BHHTEGST-Eu³⁺ & NON-EUB338-BHHTEGST-Eu³⁺ probes.

	<i>S. aureus</i>			<i>S. epidermidis</i>		
Probe	Mean Signal	Mean Background	SNR	Mean Signal	Mean Background	SNR
SAU69	95.7	3.1	31	32	6.9	4.6
EUB338	72.1	5	14.4	81	5	16.2
NONEUB338	36.7	6.1	6	33.8	7.3	4.6

Table S2: Averaged results of replicated experiments ($n_{\text{replication}}=30$) shown in Table S2: The mean signal and background staining of *S. aureus* and *S. epidermidis* are shown. The standard deviation (SDV) was calculated and is also listed.

	<i>S. aureus</i>				<i>S. epidermidis</i>			
Probe	Mean Signal	Mean Background	SNR	SDV	Mean Signal	Mean Background	SNR	SDV
SAU69	105	3	35	3.3	31	7	4.4	6.9
EUB338	75	5	15	4.6	80	5	16	5.7
NONEUB338	30	6	5	4.4	27	7	3.8	4.9