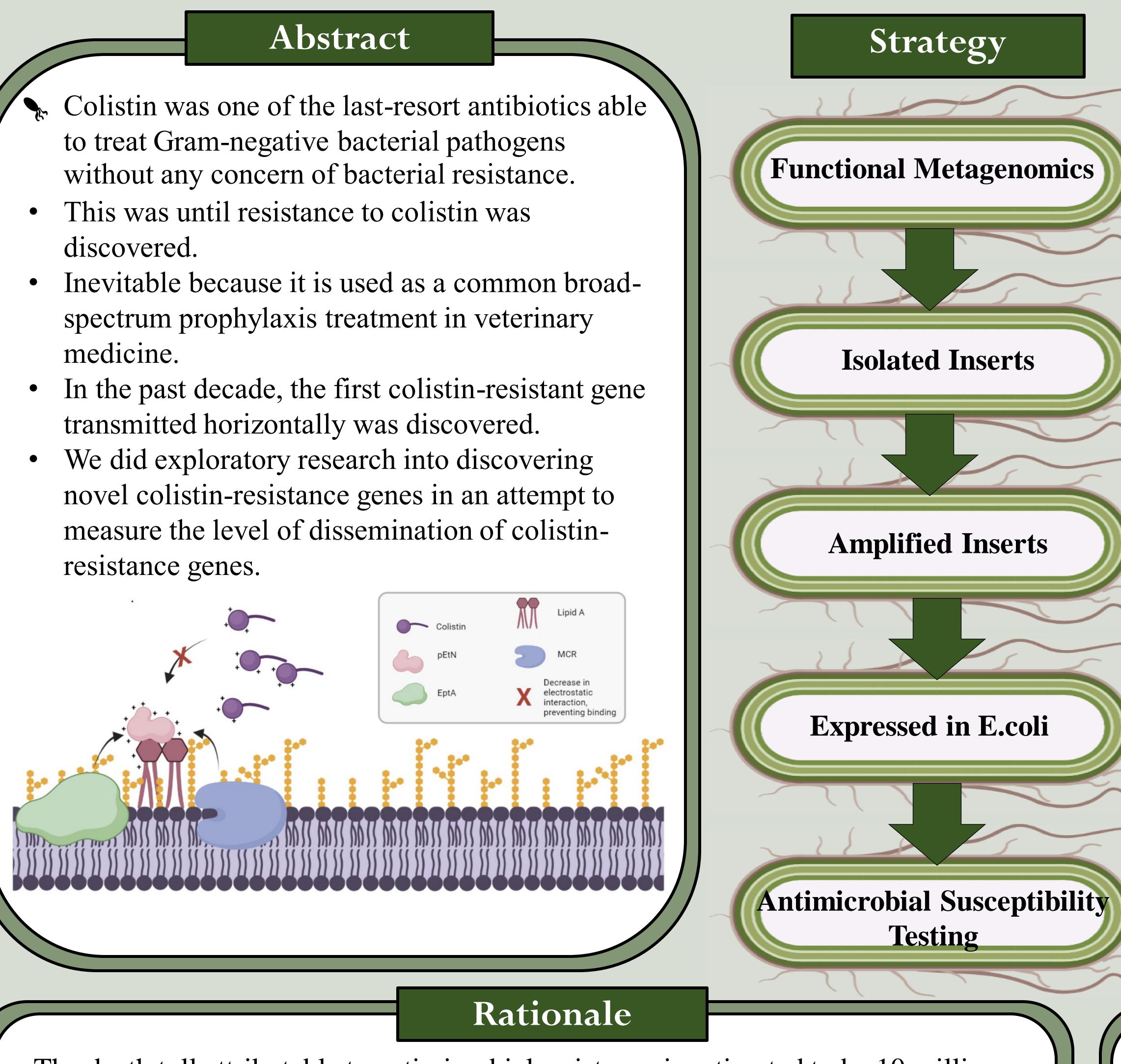


Novel Antibiotic-Resistance Genes in Goose Fecal Microbiota

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- discovered.
- medicine.
- resistance genes.



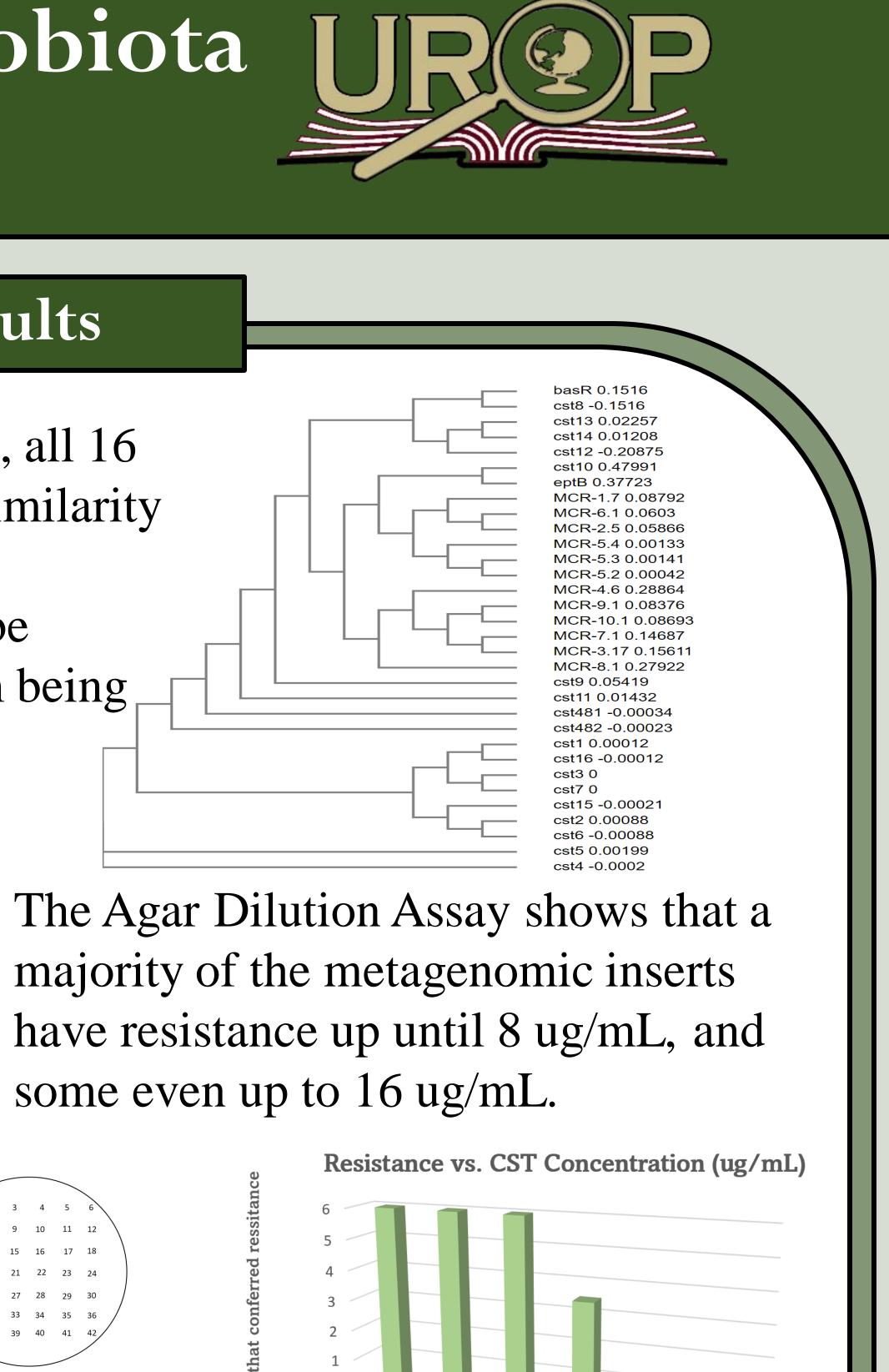
- The death toll attributable to antimicrobial resistance is estimated to be 10 million a year in 2050.
- Our findings contribute to surveillance monitoring that is vital for measuring the spread of resistance and demonstrate the consequence of prophylaxis treatment. What can be done to curb this ever-expanding issue is through acquiring knowledge of new resistance genes to potentially design antibiotics bacteria cannot resist.

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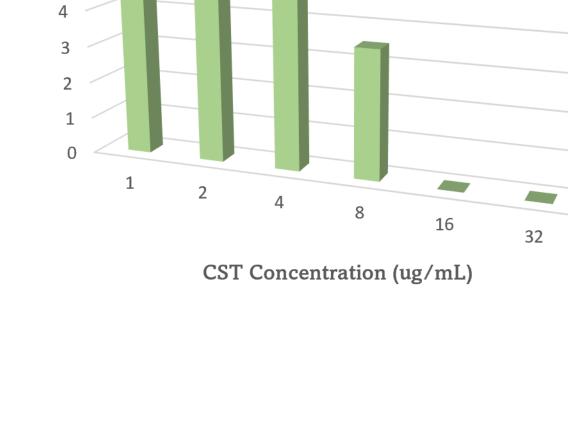
References

Results

• As shown by the phylogenetic tree, all 16 genes show significant sequence similarity to each other, MCR family, and pEtN transferases which will be verified once they come back from being_ sequenced. Concentration ug/mI 15-18 CST 6 19-22 23-26 27-28 29-32 **CST 11** 33-34 8 9 10 11 12 1, 2, 4 ug/ml 16 ug/mI 8 ug/mL The Microbroth Dilution **Future Work** Biochemical analyses The functioning capacity of microbes and microorganisms are limited in vitro. Although *E.coli* expression gives important biological information regarding the function of the gene, it does not fully reflect activity in the gene's original host. This raises the question as to whether E.coli is able to express foreign genes to their fullest potential. Further work includes protein purification for analysis of enzyme characteristics.







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|---|----|----|----|----|----|----|---|
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| | 13 | 14 | 15 | 16 | 17 | 18 | ١ |
| | 19 | 20 | 21 | 22 | 23 | 24 | |
| | 25 | 26 | 27 | 28 | 29 | 30 | / |
| | 31 | 32 | 33 | 34 | 35 | 36 | |
| | 37 | 38 | 39 | 40 | 41 | 42 | / |
| | | | | | | | |

