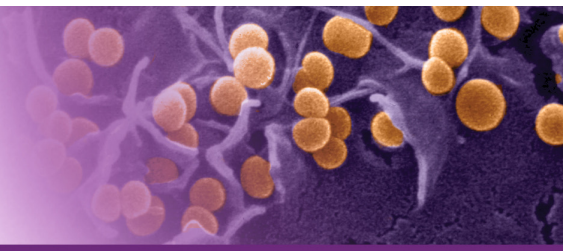


MRSA gene hunt

Gene ID cards



The activity

Using the gene ID cards you will research and classify key genes which play an important role in drug resistance and virulence for two strains of *Staphylococcus aureus* (MSSA 476 and MRSA MW2). The table (pages 3 and 4) lists 14 different genes which can be found in the genomes of the two bacterial strains, these can be found on the gene ID cards. Once you have completed the table, cluster the relevant cards into the five protein classification categories and answer the questions on page 5.

Understanding the gene ID cards

Each group has a set of 16 gene ID cards labelled with a gene name. On the back of each card is the name of the gene, a description of the protein product the gene encodes and the strain of bacteria it is found in.

hsdM, hsdS, hsdR: restriction modification system

These proteins are components that allow the bacteria to protect its DNA and recognise foreign pieces of DNA that enter the cell. This is a key defence mechanism against invasion. The hsdM modification enzyme adds chemical markers to the bacteria's own DNA. The hsdS specificity enzyme recognises these markers. However, if foreign DNA enters the cell, hsdS will recognise it as foreign, unmodified DNA, and the hsdR enzyme will chop it up.

Mobile DNA 'cassettes' or elements tend to carry this suite of enzymes. The thought is that these pieces of DNA might be protecting themselves, but by incorporating them, the bacteria benefits. Specific lineages of *S. aureus* are known to regularly host these types of defence systems; they could be ways of fit strains establishing some barriers to protect against 'free-for-all' horizontal transfer of DNA.



Staphylococcus aureus
MSSA 476



Gene ID: gene name



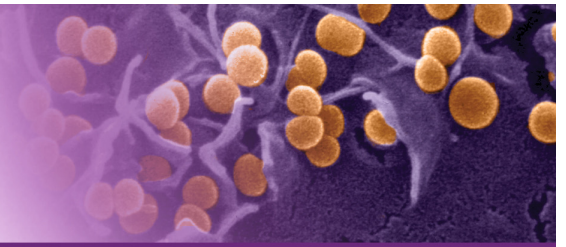
Description of the protein product and function



Strain of bacteria

MRSA gene hunt

Gene ID cards



Background information on the pathogens

MSSA 476

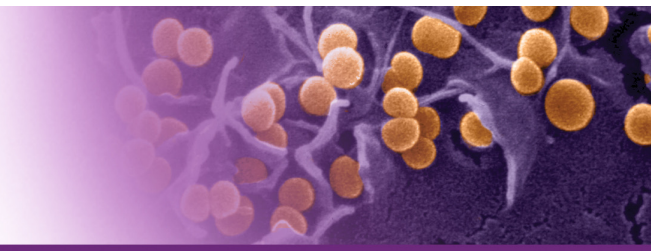
This is a methicillin-susceptible strain of *S. aureus* that was isolated from a 9-year old boy with a community-acquired infection in 1998. This strain is resistant to penicillin and fusidic acid, but can be killed by many other antibiotics. After antibiotic treatment, this patient made a full recovery. The 2,799,802 bp genome of MSSA 476 was fully sequenced by staff at the Wellcome Trust Sanger Institute in 2004.

MRSA MW2

This strain of *S. aureus* is resistant to methicillin. It was isolated in 1998 from a 16-month old girl with severe septicaemia (blood poisoning). MW2 is sensitive to some types of antibiotic, but is extremely virulent; this patient died within two hours of arriving in hospital. The genome of MW2 was fully sequenced in 2002 by staff at Juntendo University in Japan.

MRSA gene hunt

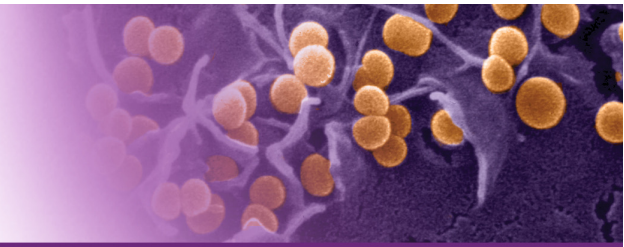
Gene ID cards



Gene	Protein product	Role/Function	MSSA 476	MRSA MW2	Classification
<i>ccrA & ccrB</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>ear</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>int</i> (region 3)					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>fusB1</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>hsdR, hsdS, hsdM</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>

MRSA gene hunt

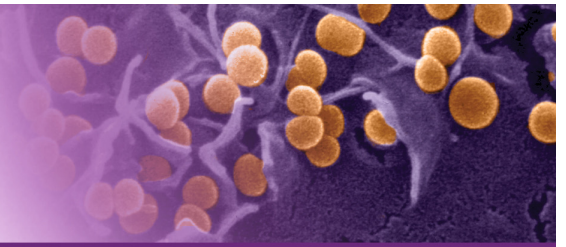
Gene ID cards



Gene	Protein product	Role/Function	MSSA 476	MRSA MW2	Classification
<i>orfX</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>lukS & lukF</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>mecA</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>sec4</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>
<i>sel2</i>					Antibiotic resistance <input type="checkbox"/> Enzyme <input type="checkbox"/> Surface protein <input type="checkbox"/> Toxin <input type="checkbox"/> Mobile genetic element <input type="checkbox"/>

MRSA gene hunt

Gene ID cards



Using the flipchart paper, classify the genes into five categories: antibiotic resistance, enzyme, surface protein, toxin, mobile genetic element. Stick the cards under their correct corresponding headings. Feed back the results to the class and be prepared to discuss the following questions:

Q1a) Which genes are found in MSSA 476?

Q1b) Which genes are found in MRSA MW2?

Q1c) Which genes are found in both strains?

Q2. Which genes could be termed as virulence factors, i.e. factors that give the bacterium a greater capacity to cause disease?

Q3. Which genes might give one strain an advantage over other strains?

Q4. What processes do you think underlie the evolution and spread of antibiotic-resistance in bacteria?