Emergence of a novel strain of healthcare associated MRSA

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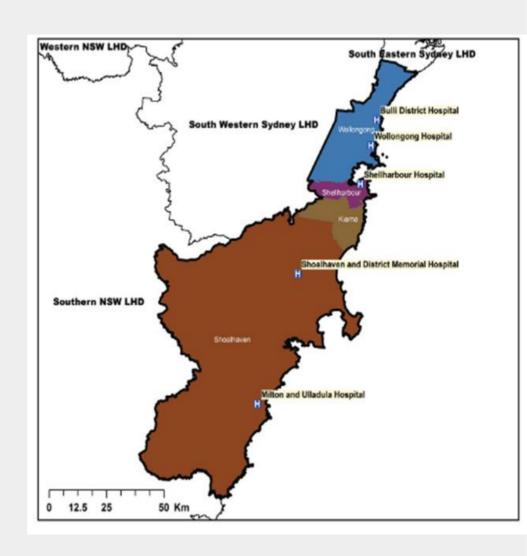
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ISLHD Infection Management and Control Service



Illawarra Shoalhaven LHD

- Between the mountains and the sea
- Population of 368,822 is projected to reach over 425,000 by 2021
- 9 inpatient hospitals and numerous community-based services



IMACS' routine surveillance for multiresistant organisms (MROs)

- Daily download from SEALS for all MROs
- IMACS CNC assesses MRSA antibiotic resistance pattern of each isolate to inform classification as H-MRSA, C-MRSA or UK-MRSA
- Attribution of likely acquisition (healthcare associated hospital/unit, or community associated)
- Entry into local MRO database to allow analysis



MRSA classification by phenotype

Antimicrobial	C-MRSA	UK-MRSA	H-MRSA	novel
			R to 2 or more Gent, Erythromycin, Tetracycline	R to 2 or more Gent, Erythromycin, Tetracycline
Methicillin	R	R	R	R
Gentamicin	S	S	R	R
Co-trimoxazole	S	S	R	S
Ciprofloxacin	S	R		

Single occurrences August and September 2013 - ?anomaly

First cluster noticed affecting 6 patients from 4 different hospitals providing rehabilitation and aged care services in November 2013

Non-invasive infections / colonisation



Local control measures

- Continued passive surveillance
- March 2014 Binary typing of all new MRSA isolates
 - Isolates described as binary type 1296
 - Reference lab confirm not previously seen in NSW
- Improved infection control precautions
 - Hand hygiene
 - Equipment cleaning
 - Environmental cleaning



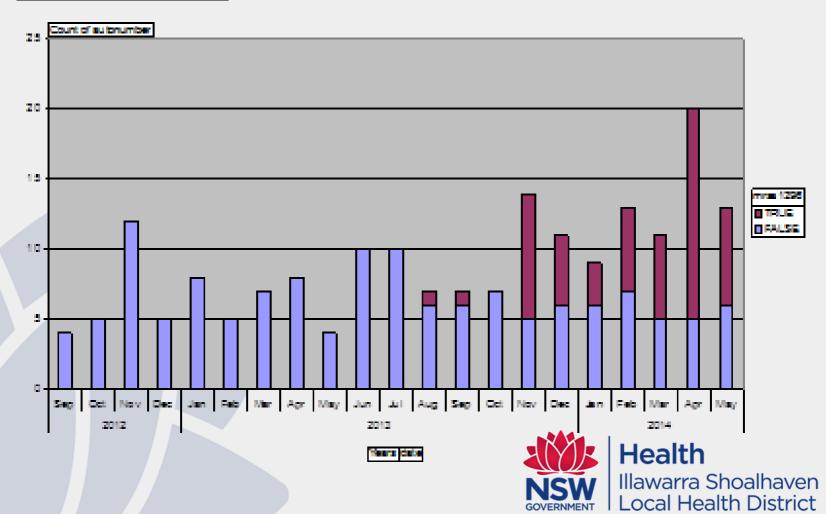
May 2014

- First bloodstream infection caused by novel MRSA
- Monitoring indicated that bt1296 was adding to rather than replacing the usual H-MRSA, and numbers were increasing
- Informal networking revealed no other LHDs noticing this phenotype
- Outbreak management committee formed



Graph May 2014

putbreak (All) has HCA species MRSAHA



Outbreak management

- Case definition
 - S. aureus, resistant to 2 or more of Gentamicin,
 Erythromycin, Tetracycline <u>plus</u> sensitive to Co-trimoxazole
- Risk Register
 - Patient outcomes
 - Increased burden of MRSA within NSW Health system
- Escalation to NSW Health and CEC
 - Expert Advisory Group established
 - Support to epidemiological study



NSW Health

- Notification to all LHDs and private hospitals with request for similar isolates to be binary typed
- Admission screening of patients transferred from ISLHD advised
- Close monitoring of situation through regular teleconferences between LHD, CEC and NSW Health



Local actions

- Presentation to executive management teams and lead clinicians for affected services
- Grand Round presentations
- Active surveillance MRSA admission and discharge screening
- Investigation of hypothesis that patient transit vehicles implicated in transmission
- Reinforcement of ISLHD 'Bare Below Elbows' policy
- Continue to ensure ABHR "6 inches from the patient's right foot"
- Successful project to improve hand hygiene led by junior doctors

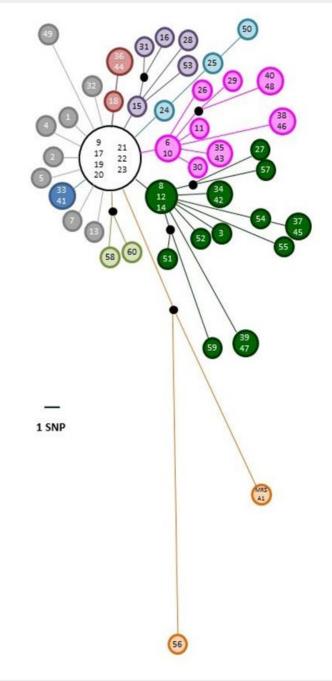
Health

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- Improved access to detergent wipes
- Approval for detailed genotyping of first isolates

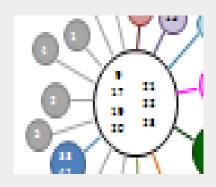
Whole genome sequencing

- 60 isolates
- Eight different clades
- Isolates in each clade have no more than 12 single nucleotide polymorphism (SNP) differences between them
- Isolates within a single circle of a clade had indistinguishable SNP profiles



White clade

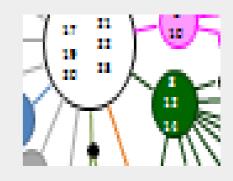
- December 2013 April 2014
- Seven patients
- Two different hospitals, three different wards, five different teams, three different specialties
- Two patients shared the same medical officer (isolates 20 and 22)
- No obvious links between other patients





Green clade

- 3 patients
- Same hospital, three different wards
- All geriatrics, three different consultants

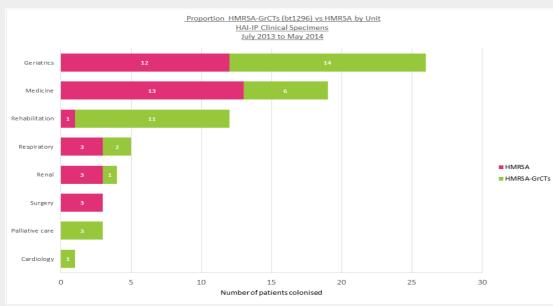




Case control comparison (67 cases;112 controls)

- Controls = patients consecutively listed on the LHD MRO database with isolates of non-novel H-MRSA strains over the same time period.
- Rate of colonisation and infection comparable
- MRSA bt1296 more likely to affect people admitted to more than one hospital in previous 90 days
- People admitted to a geriatric ward were 1.86 (1-3.43)
 times more likely to have bt1296 (p=0.05)
- 7 of the 11 deaths in people with MRSA bt1296 were in colonised people, and due to co-morbidities rather than bt1296.

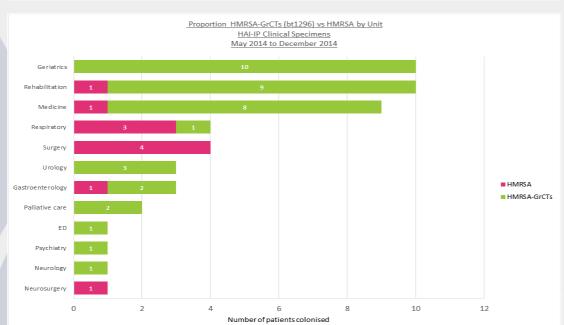
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Proportion of bt1296 by Specialty

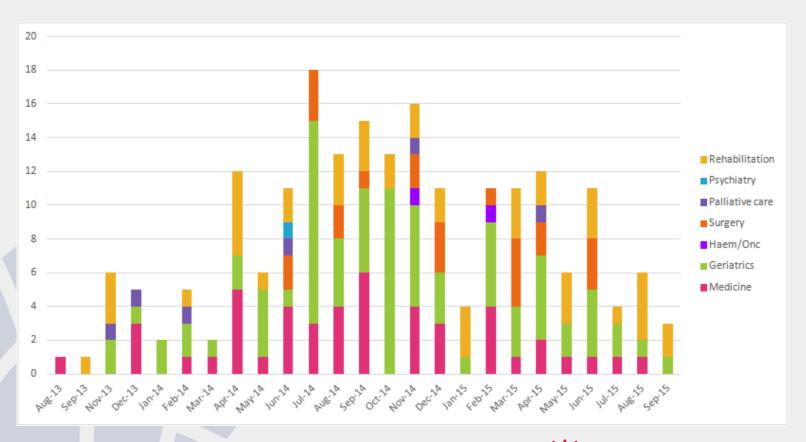
July 2013 to May 2104



June 2014 to December 2014

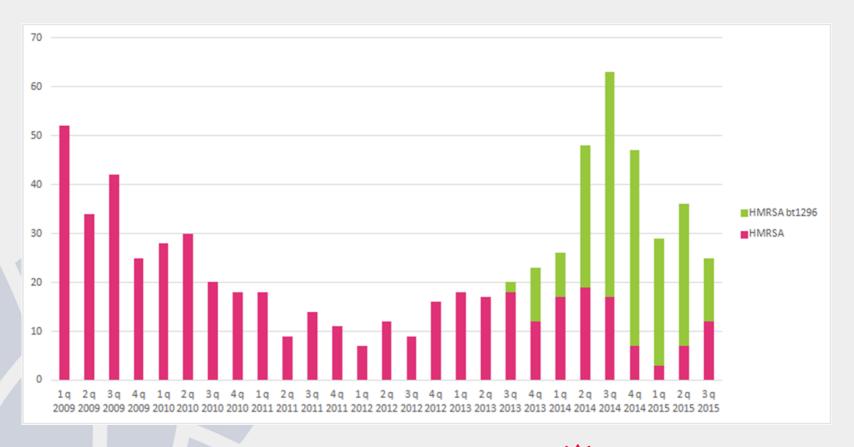


HCA-IP HMRSA bt1296 by Speciality





HCA-IP All Specimens January 2013 to September 2015





Conclusions

- Antibiogram review identified the emergence of a novel strain of MRSA
- Antibiogram correlated well with binary typing results
- Whole genome sequencing may be beneficial if results are available soon after the onset of the outbreak
- Outbreak control is extremely challenging when more than one clinical area is implicated
- Make it easy for staff to 'do the right thing'
- Effective communication can inspire others to act



Acknowledgements

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