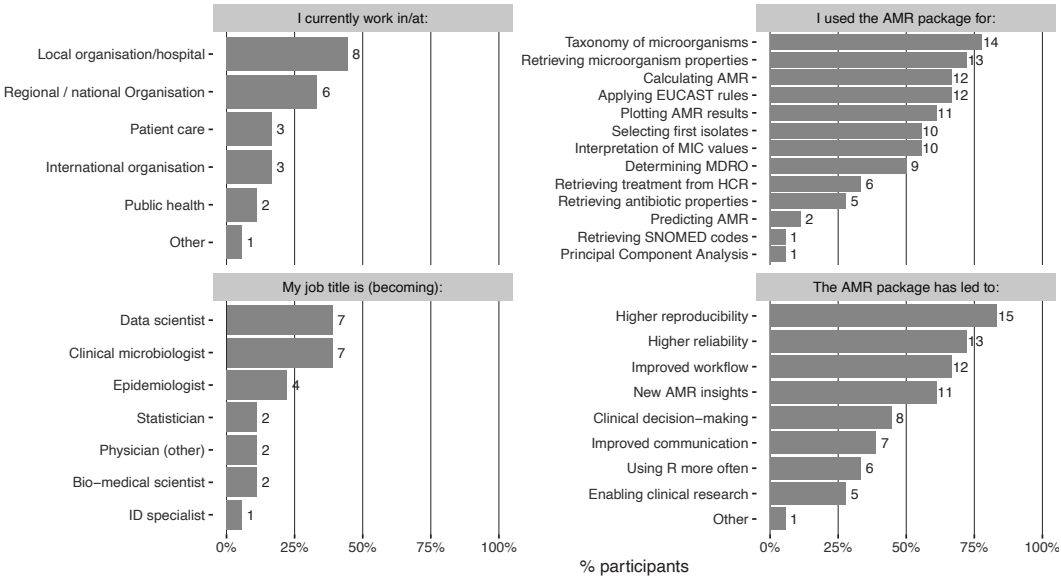
I currently work in/at:
Local organisation/hospital
8
Regional / national Organisation
6
Patient care
3
International organisation
3
Public health
2
Other
1
I used the AMR package for:
Taxonomy of microorganisms
14
Retrieving microorganism properties
13
Calculating AMR
12
Applying EUCAST rules
12
Plotting AMR results
11
Selecting first isolates
10
Interpretation of MIC values
10
Determining MDRO
9
Retrieving treatment from HCR
6
Retrieving antibiotic properties
5
Predicting AMR
2
Retrieving SNOMED codes
1
Principal Component Analysis
1
My job title is (becoming):
Data scientist
7
Clinical microbiologist
7
Epidemiologist
4
Statistician
2
Physician (other)
2
Bio−medical scientist
2
ID specialist
1
The AMR package has led to:
Higher reproducibility
15
Higher reliability
13
Improved workflow
12
New AMR insights
11
Clinical decision−making
8
Improved communication
7
Using R more often
6
Enabling clinical research
5
Other
1
0%
25%
50%
75%
100%
% participants