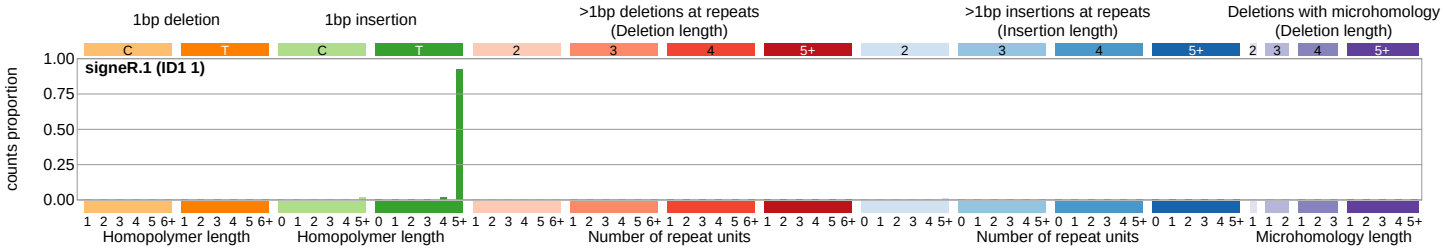
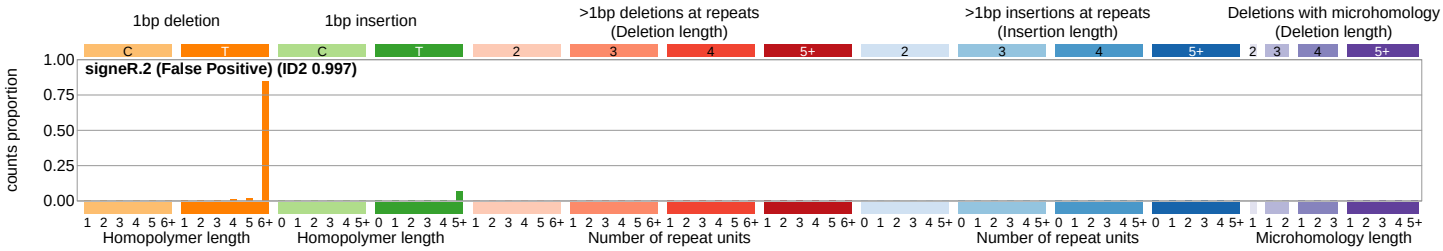

1bp deletion
1bp insertion
>1bp deletions at repeats (Deletion length)
>1bp insertions at repeats (Insertion length)
Deletions with microhomology (Deletion length)
signeR.1 (ID1 1)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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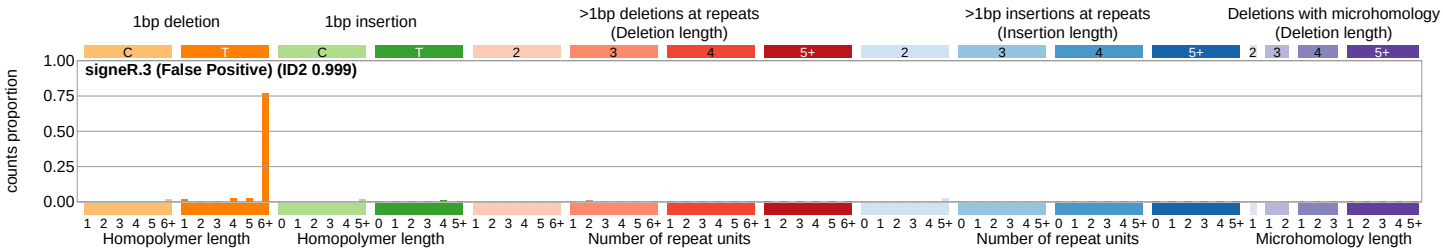
Deletions with microhomology (Deletion length)
signeR.2 (False Positive) (ID2 0.997)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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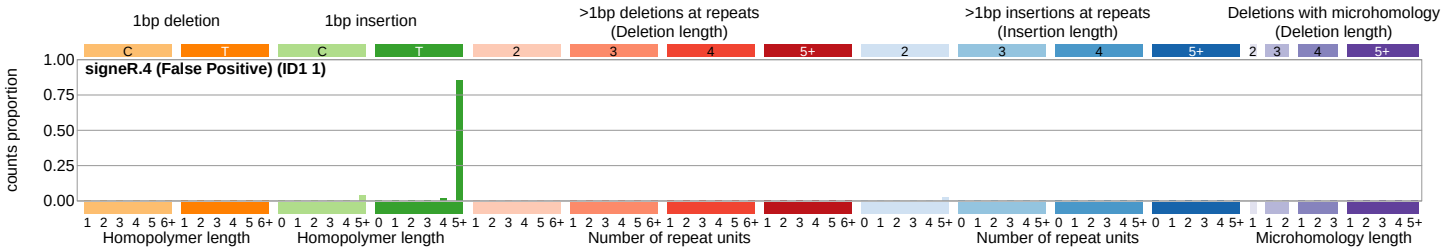
Deletions with microhomology (Deletion length)
signeR.3 (False Positive) (ID2 0.999)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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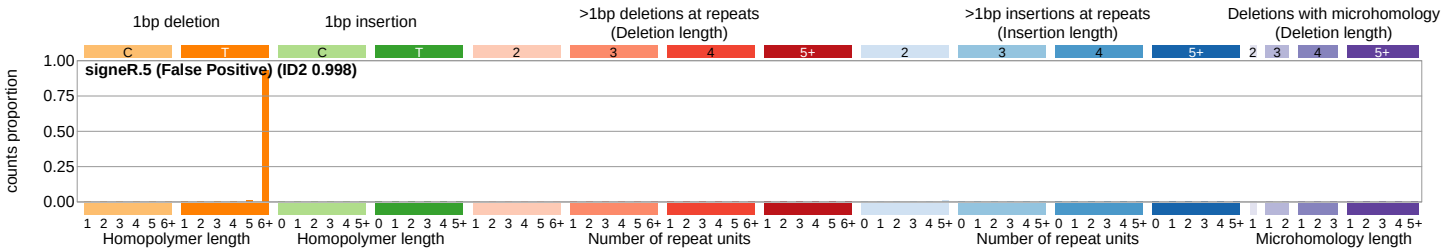
Deletions with microhomology (Deletion length)
signeR.4 (False Positive) (ID1 1)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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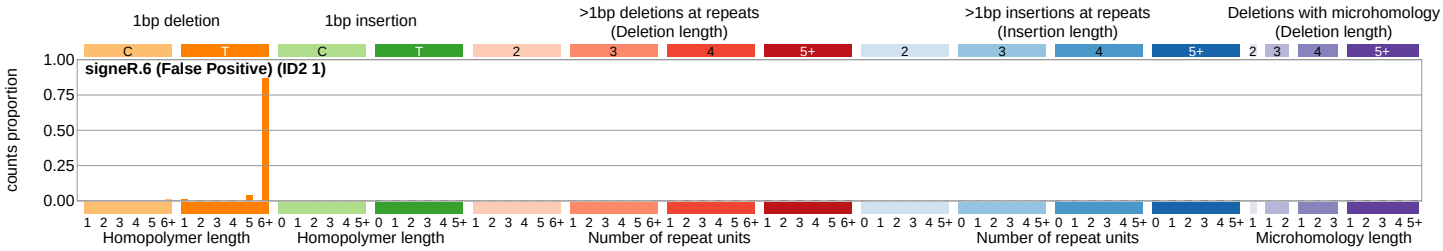
Deletions with microhomology (Deletion length)
signeR.5 (False Positive) (ID2 0.998)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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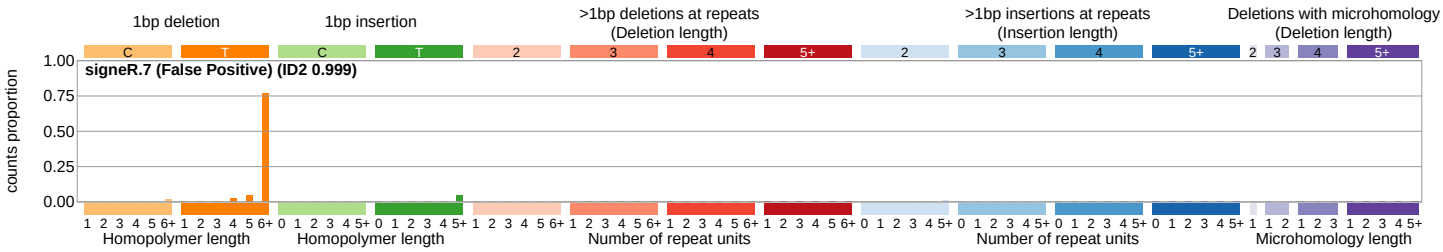
Deletions with microhomology (Deletion length)
signeR.6 (False Positive) (ID2 1)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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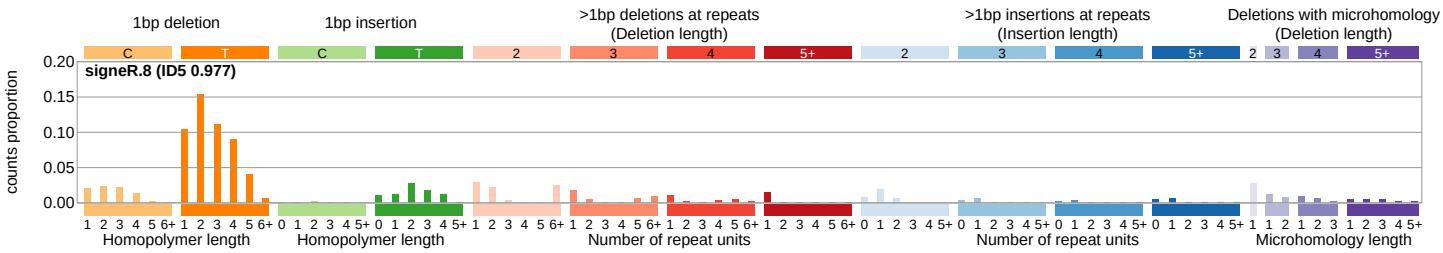
Deletions with microhomology (Deletion length)
signeR.7 (False Positive) (ID2 0.999)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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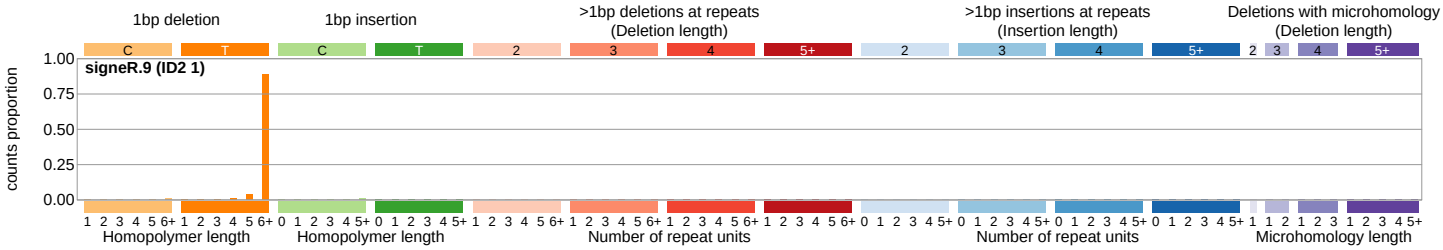
Deletions with microhomology (Deletion length)
signeR.8 (ID5 0.977)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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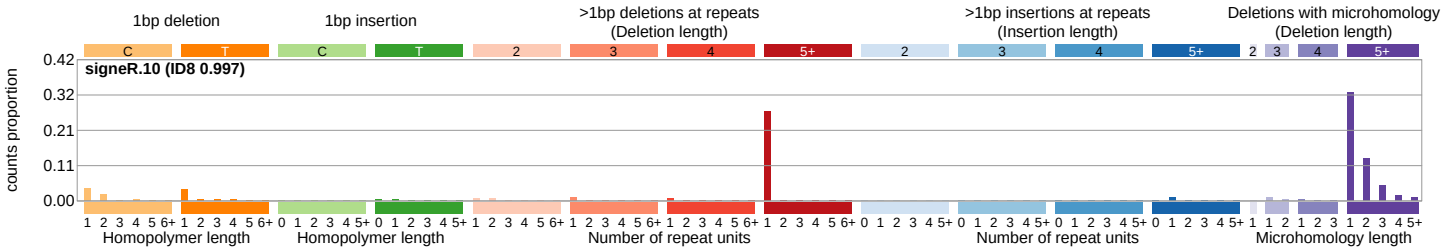
Deletions with microhomology (Deletion length)
signeR.9 (ID2 1)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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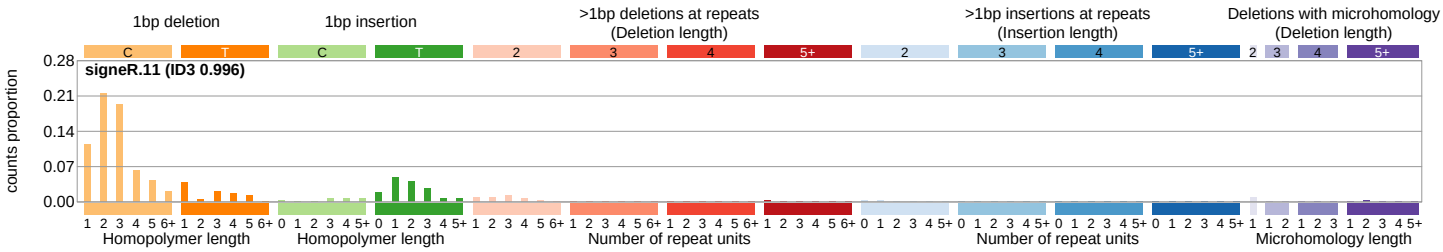
Deletions with microhomology (Deletion length)
signeR.10 (ID8 0.997)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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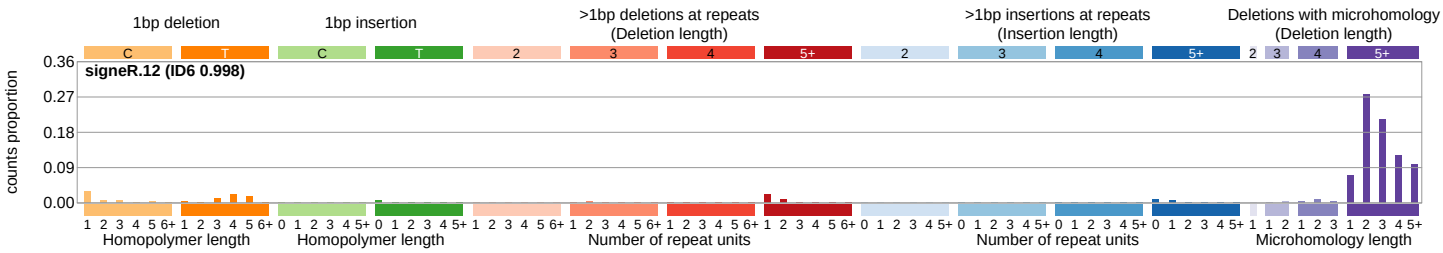
Deletions with microhomology (Deletion length)
signeR.11 (ID3 0.996)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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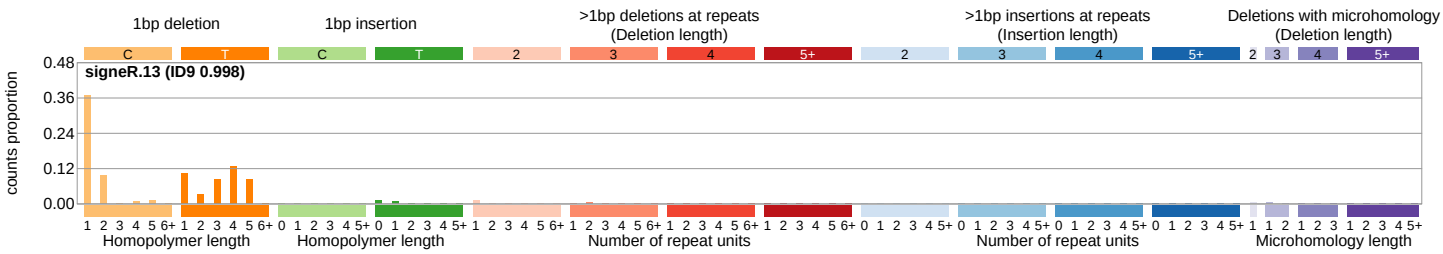
Deletions with microhomology (Deletion length)
signeR.12 (ID6 0.998)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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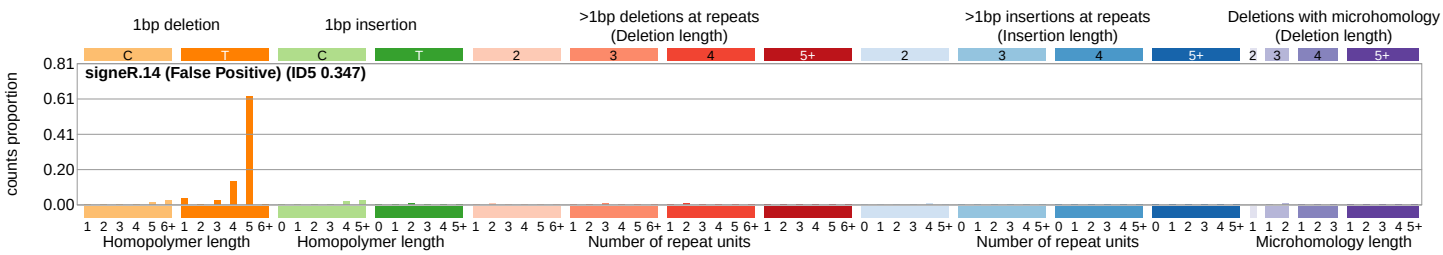
Deletions with microhomology (Deletion length)
signeR.13 (ID9 0.998)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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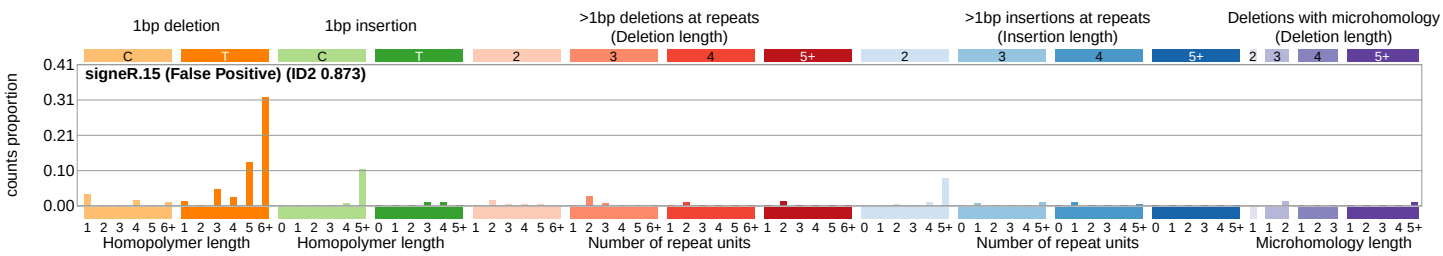
Deletions with microhomology (Deletion length)
signeR.14 (False Positive) (ID5 0.347)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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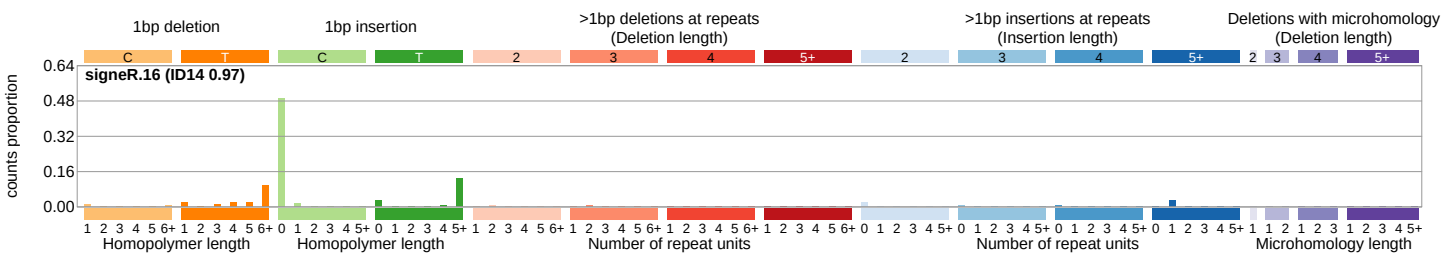
Deletions with microhomology (Deletion length)
signeR.15 (False Positive) (ID2 0.873)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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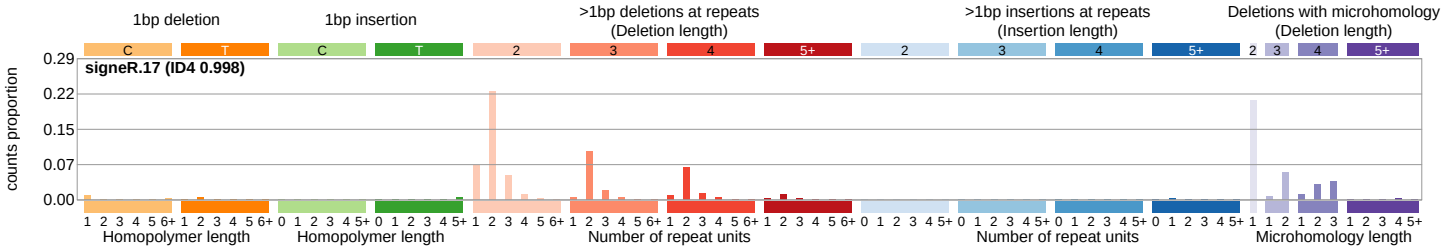
Deletions with microhomology (Deletion length)
signeR.16 (ID14 0.97)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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1bp insertion
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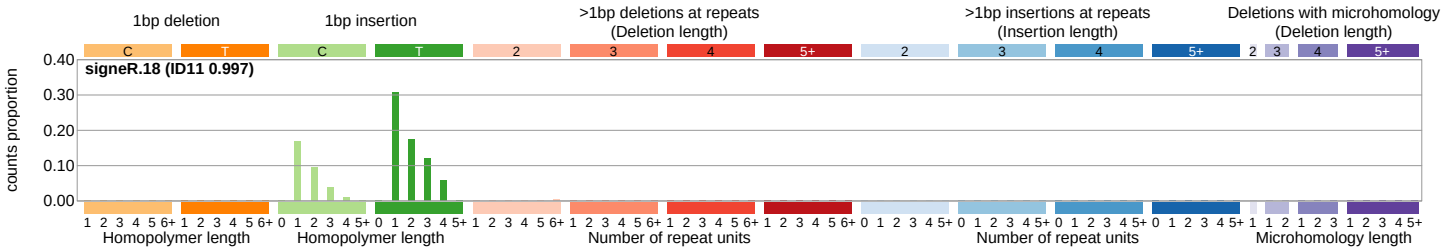
>1bp insertions at repeats (Insertion length)
Deletions with microhomology (Deletion length)
signeR.17 (ID4 0.998)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length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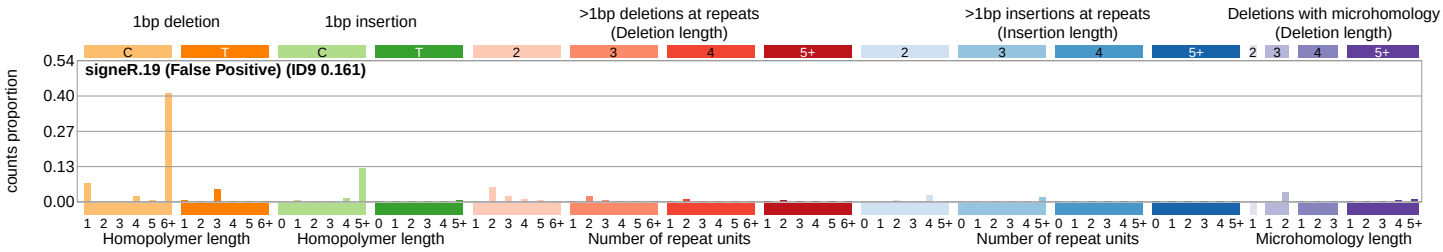
1bp insertion
>1bp deletions at repeats (Deletion length)
>1bp insertions at repeats (Insertion length)
Deletions with microhomology (Deletion length)
signeR.18 (ID11 0.997)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length


1bp deletion
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Deletions with microhomology (Deletion length)
signeR.19 (False Positive) (ID9 0.161)
counts proportion
Homopolymer length
Number of repeat units
Microhomology length