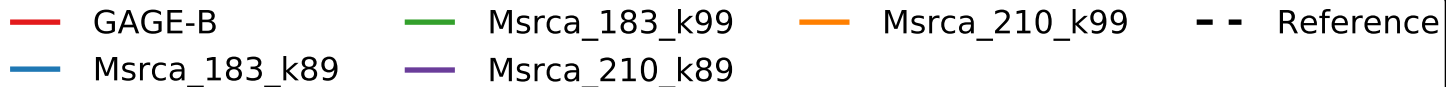
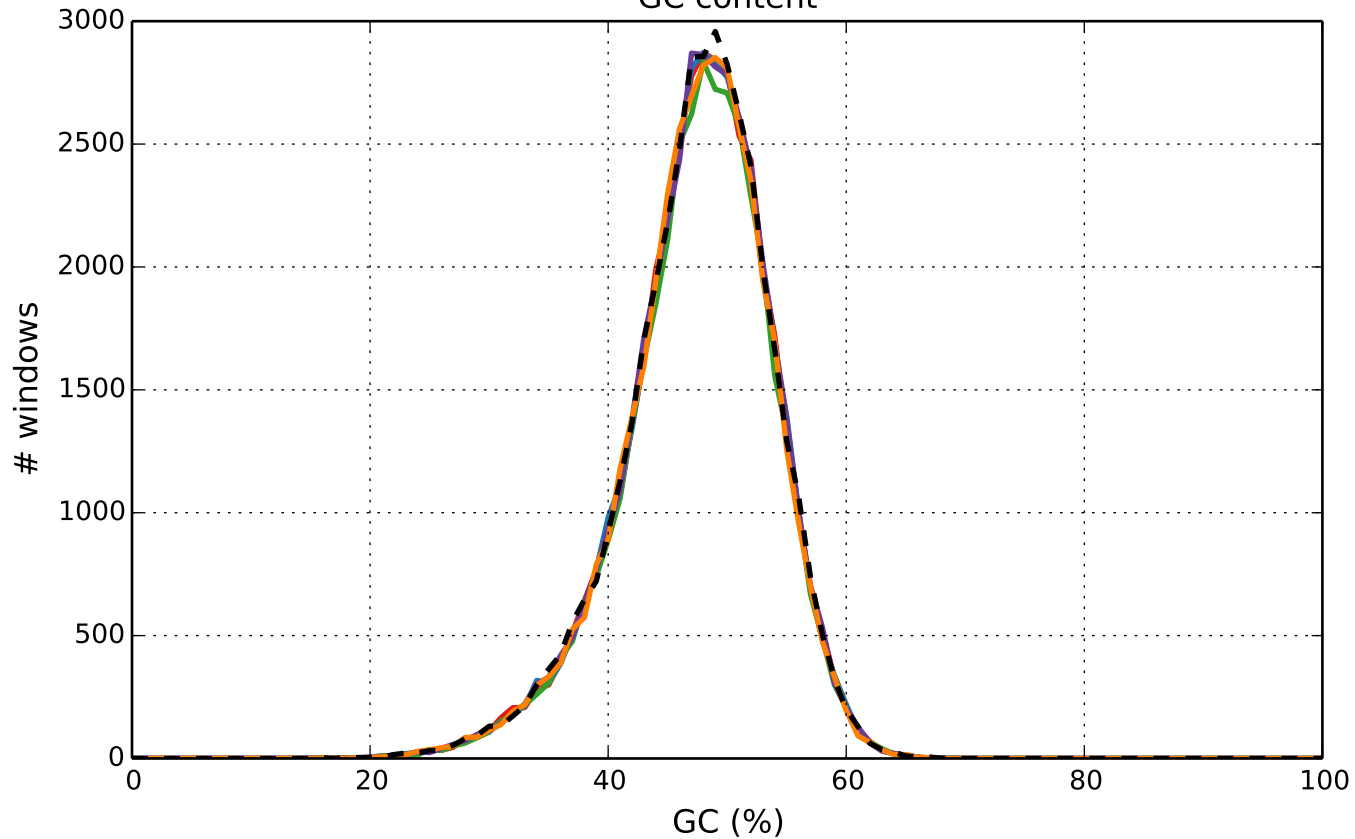
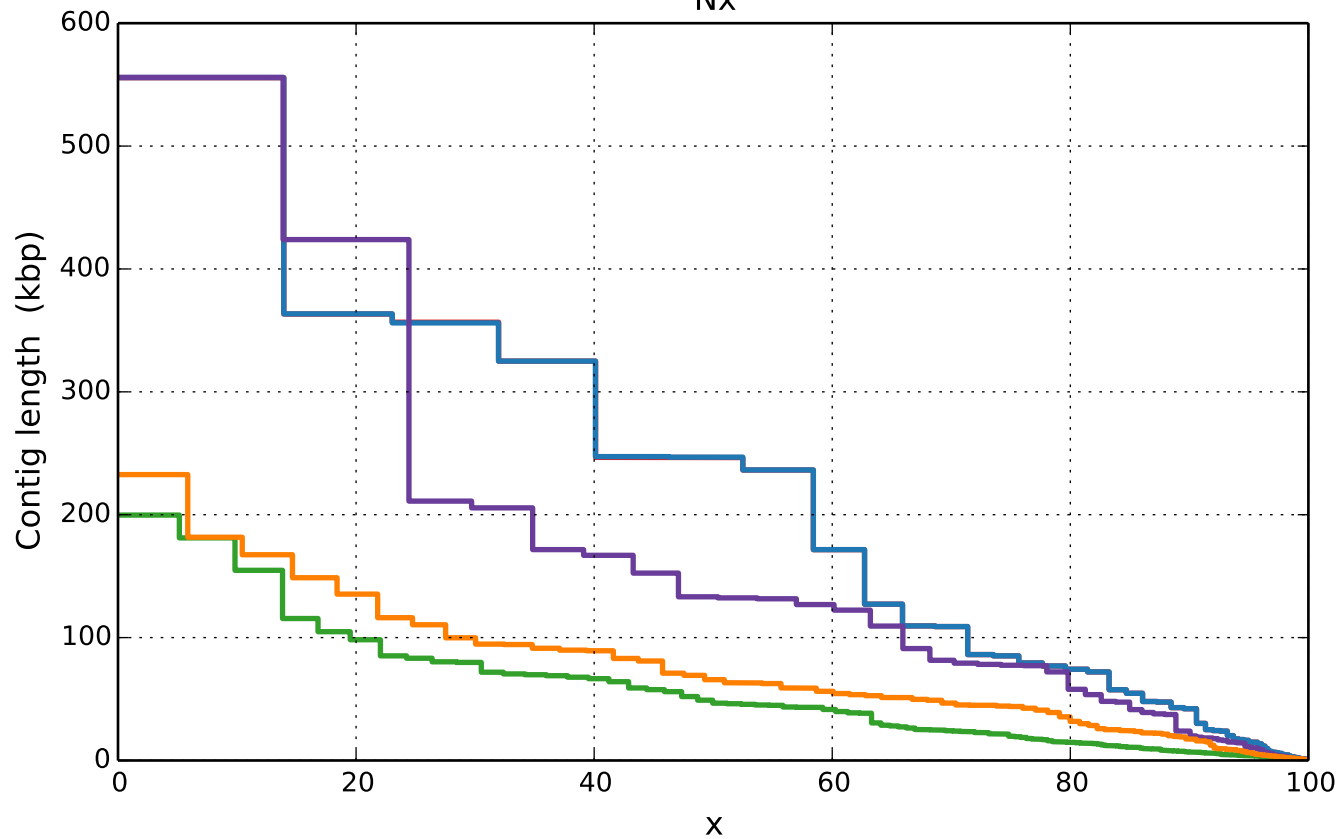


GC content



Nx



GAGE-B

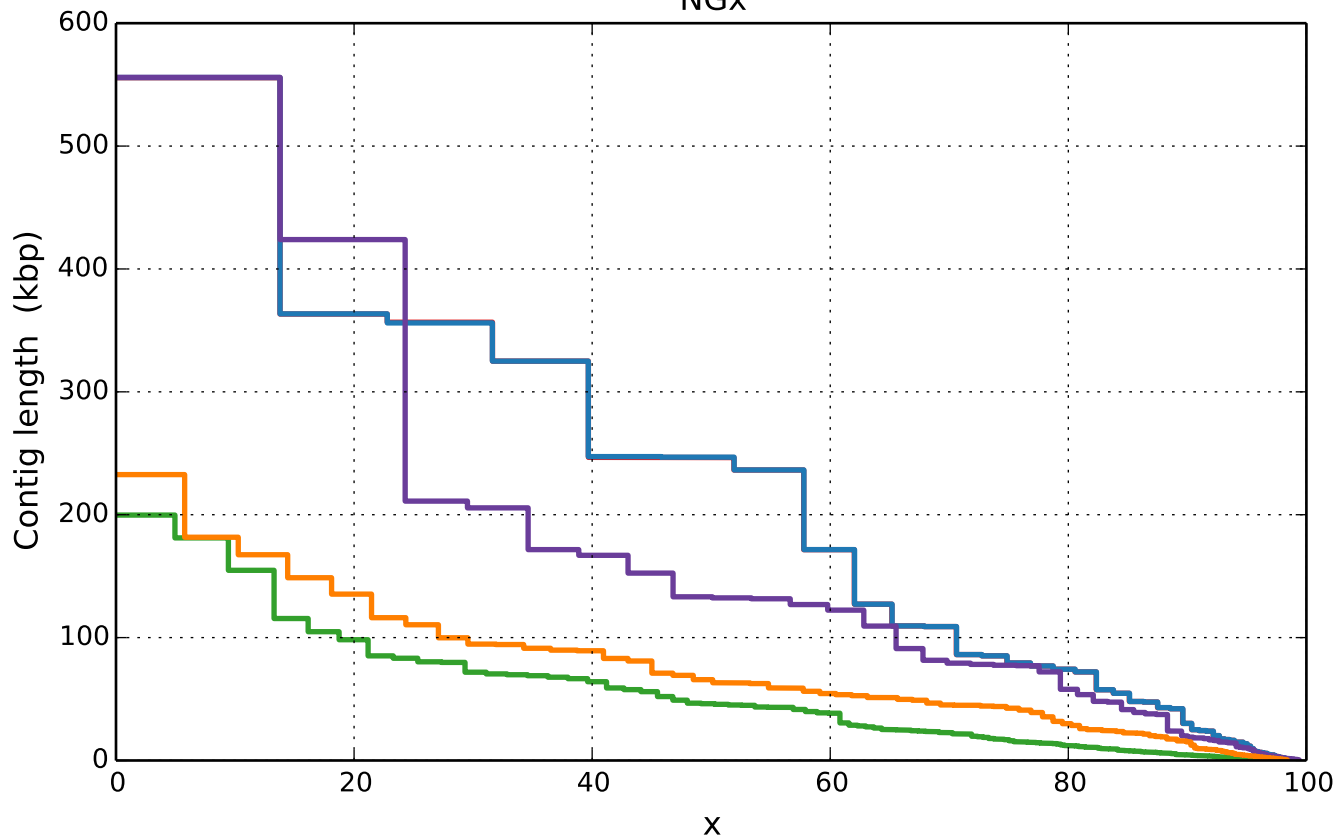
Msrca\_183\_k99

Msrca\_210\_k89

Msrca\_210\_k99

Msrca\_183\_k89

NGx



GAGE-B

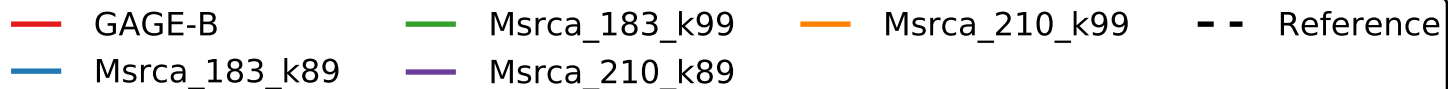
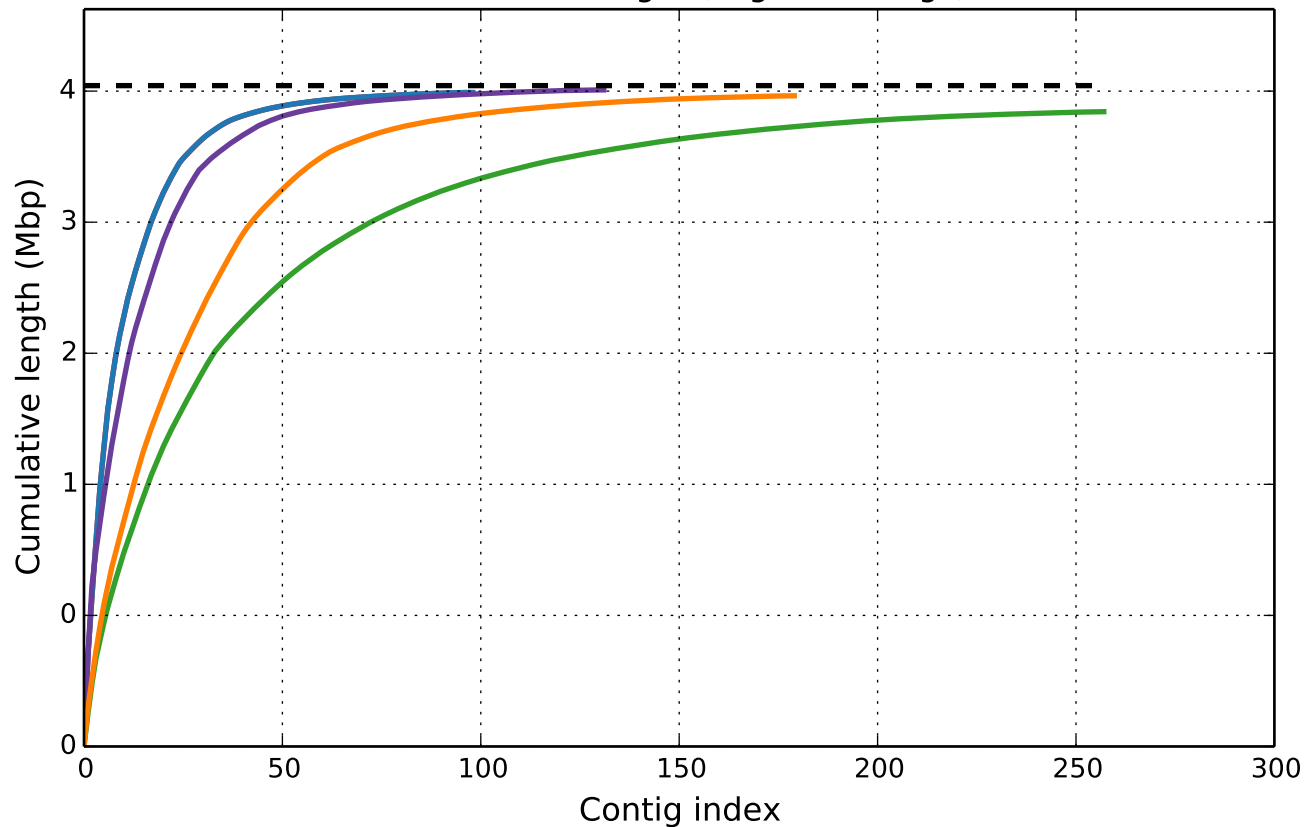
Msrca\_183\_k99

Msrca\_210\_k89

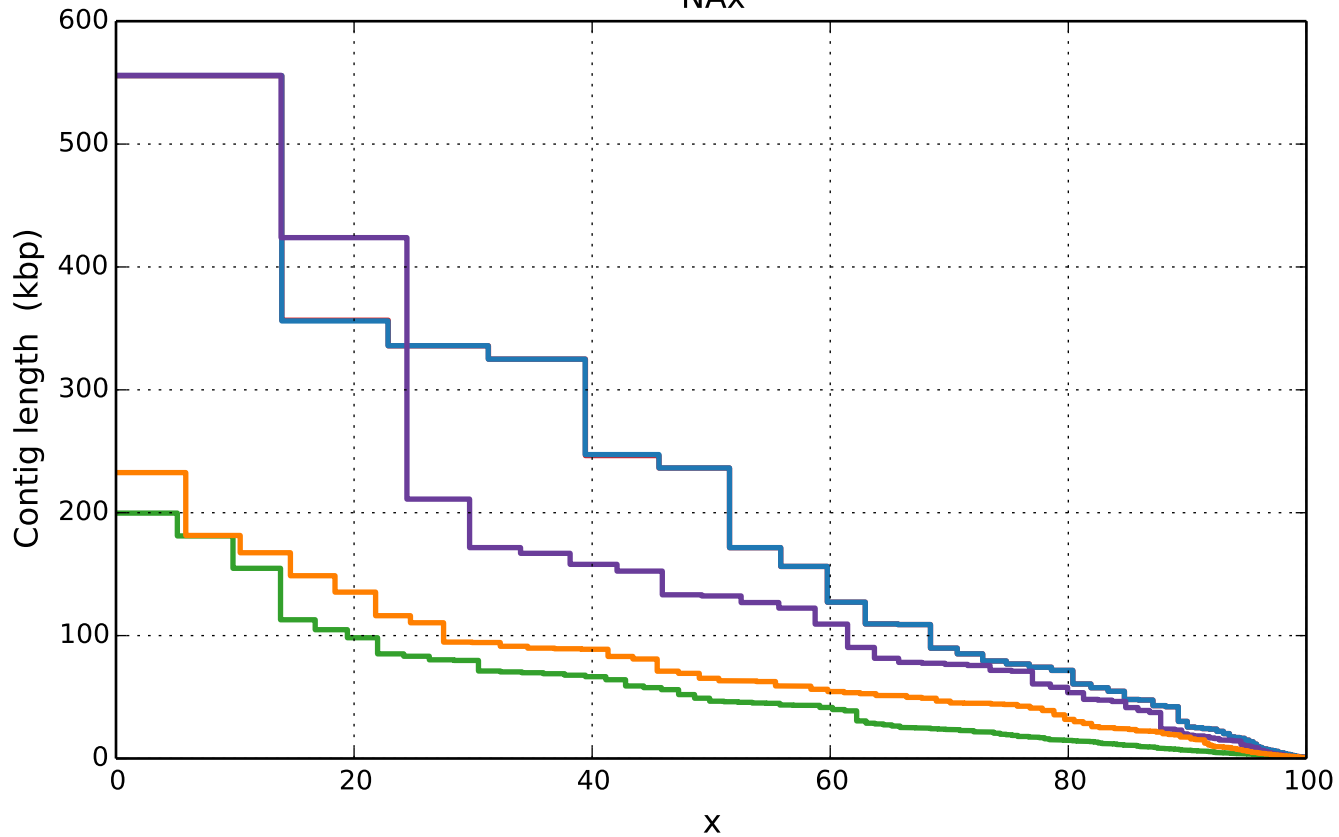
Msrca\_210\_k99

Msrca\_183\_k89

Cumulative length (aligned contigs)



NAx



GAGE-B

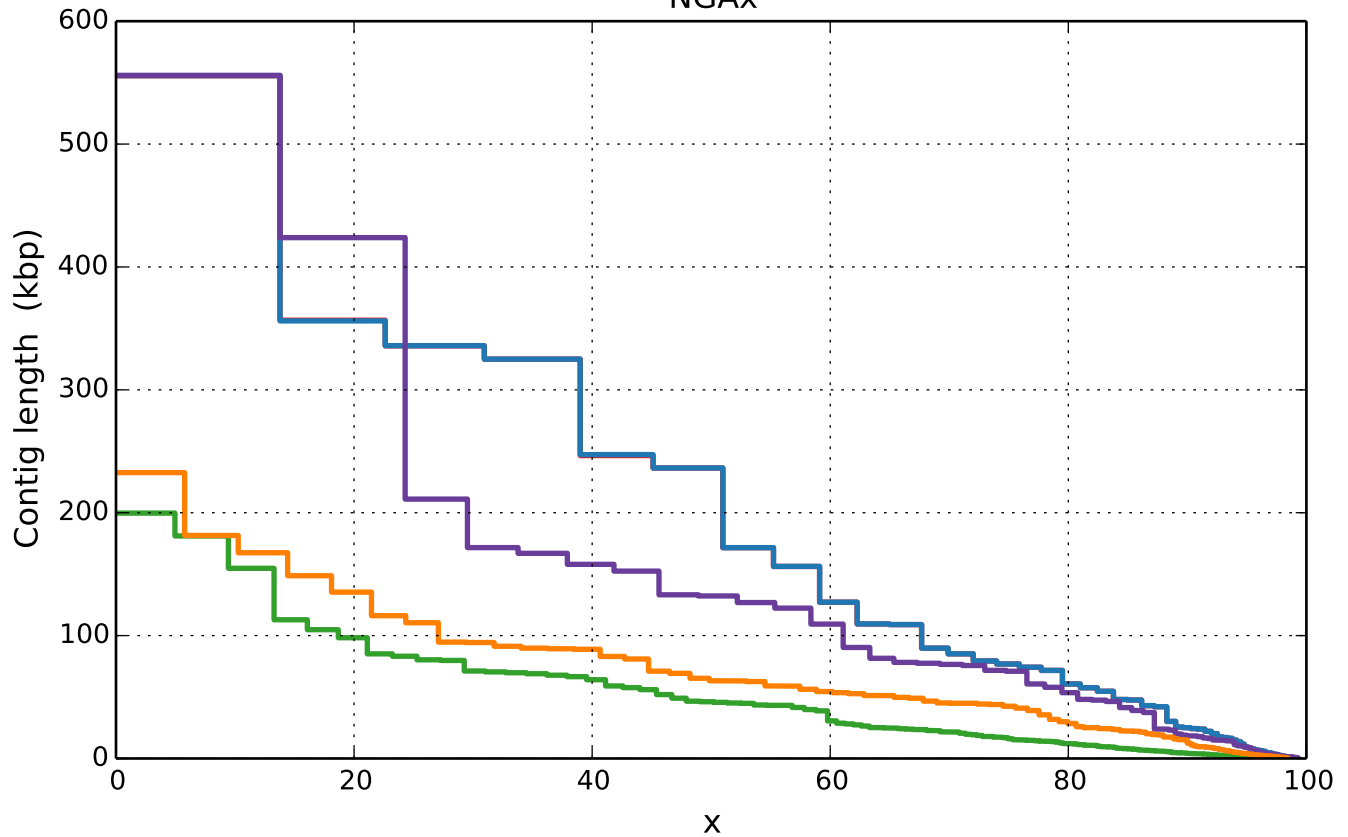
Msrca\_183\_k99

Msrca\_210\_k89

Msrca\_210\_k99

Msrca\_183\_k89

NGAx



GAGE-B

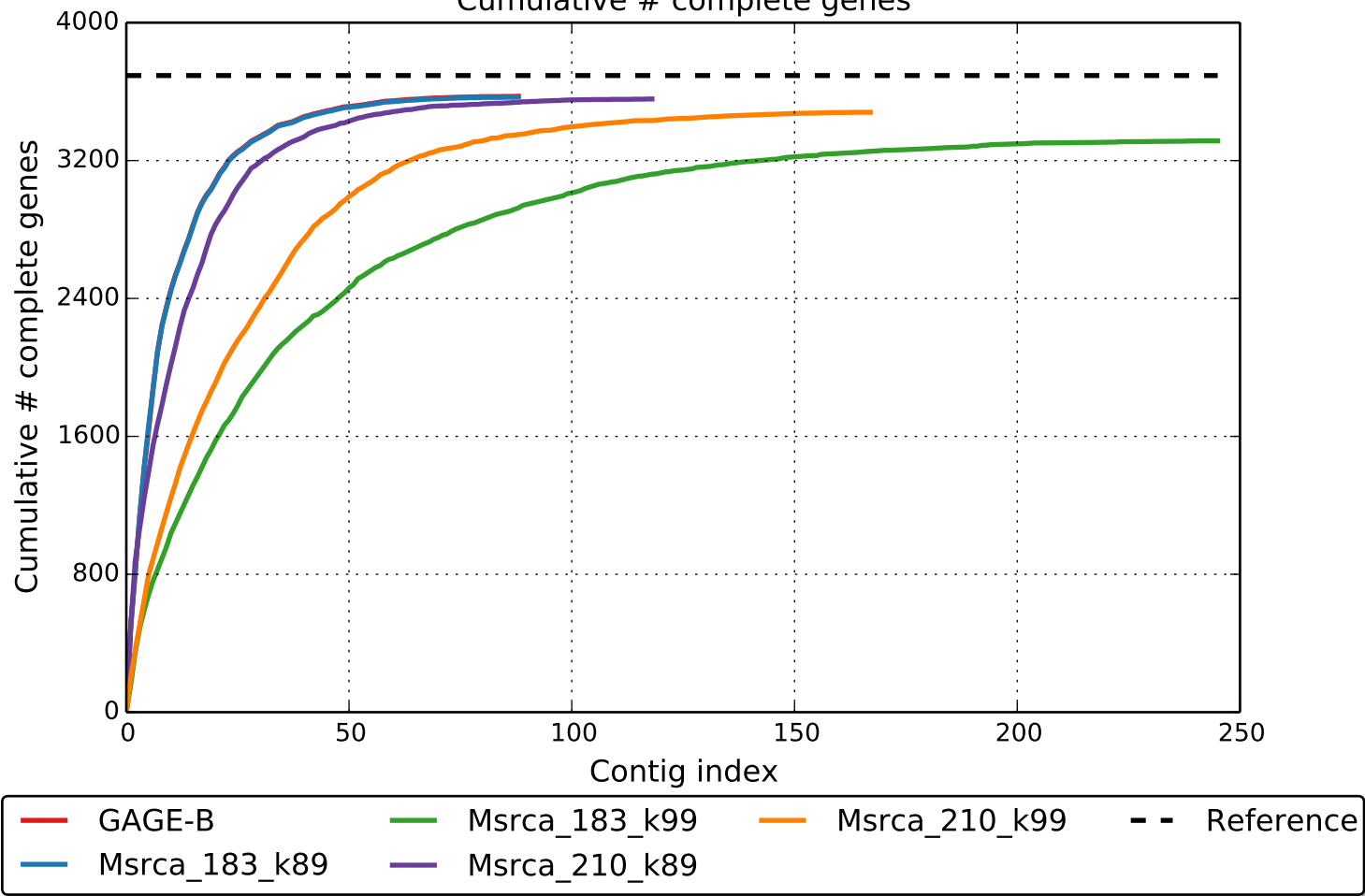
Msrca\_183\_k99

Msrca\_210\_k89

Msrca\_210\_k99

Msrca\_183\_k89

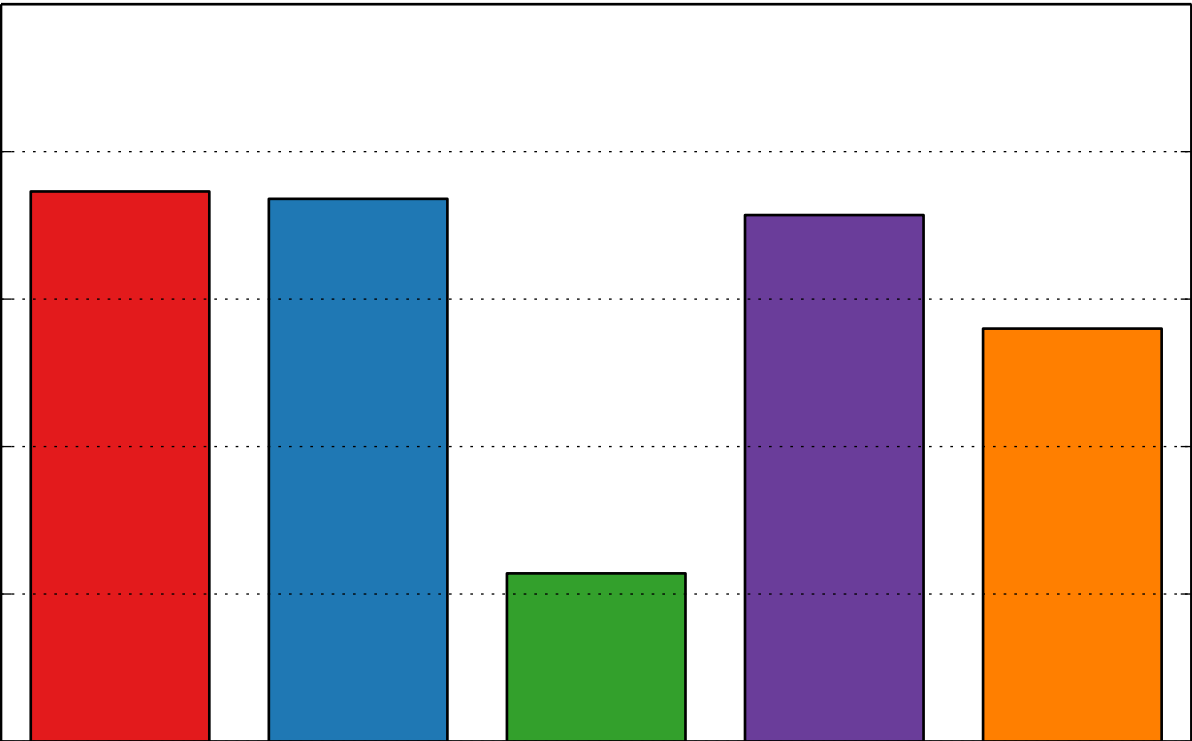
Cumulative # complete genes





# complete genes

3700  
3600  
3500  
3400  
3300  
3200



GAGE-B

Msrca\_183\_k99

Msrca\_210\_k89

Msrca\_210\_k99

Msrca\_183\_k89

