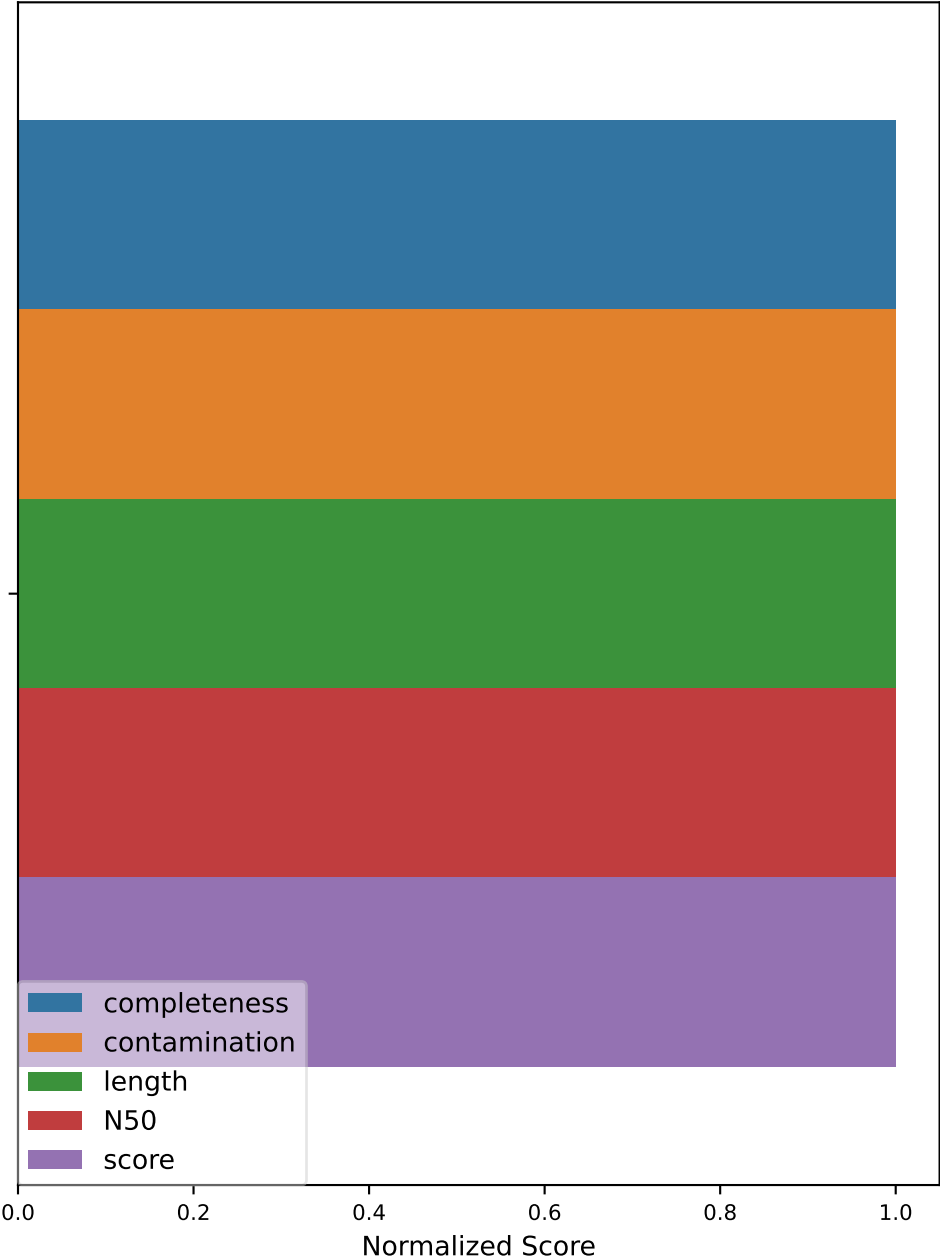


Scoring of cluster 1\_0

genome

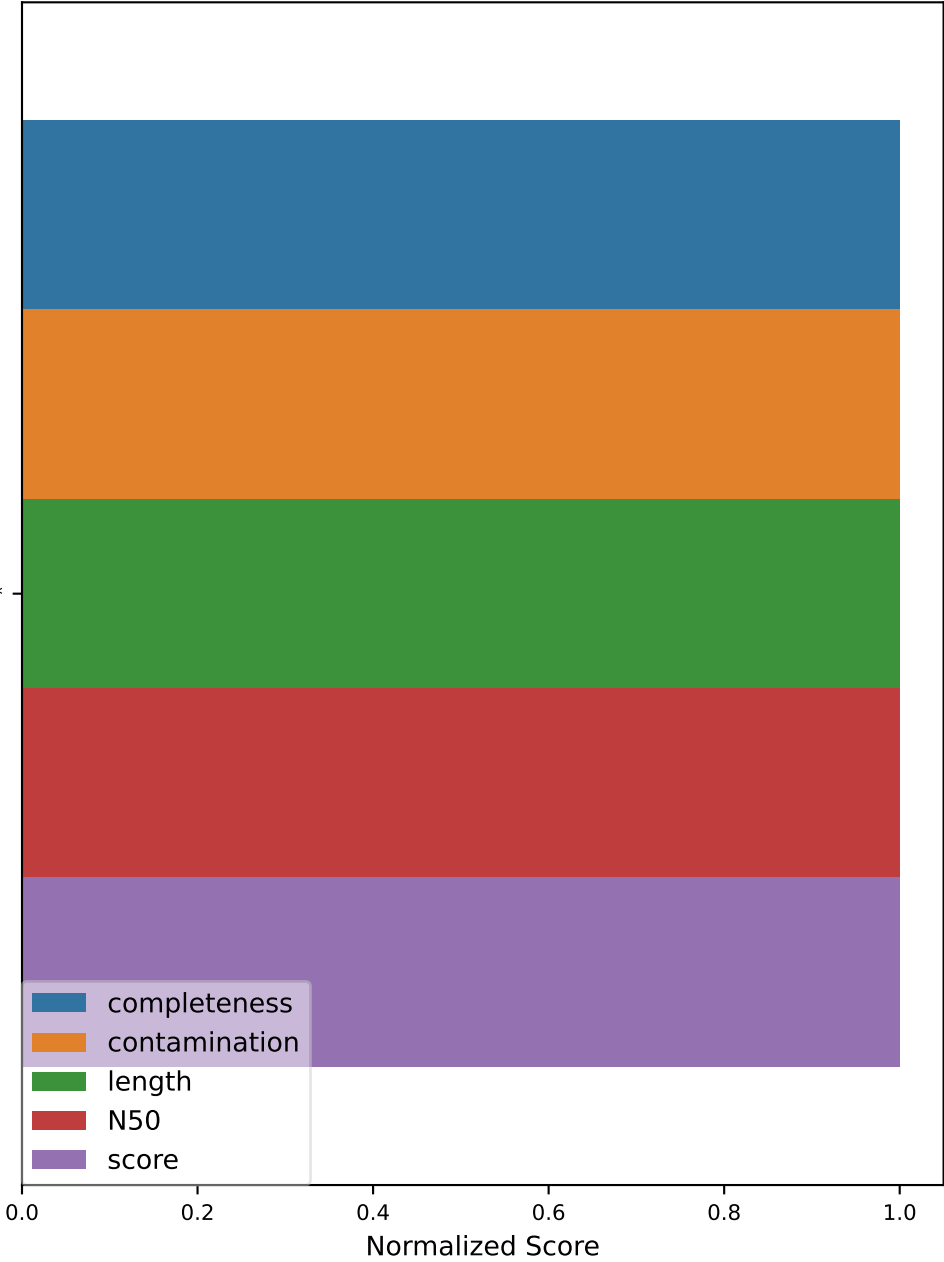
49647\_1#106\_clean\_bin.23\_34\_filtered\_kept\_contigs.fasta.gz \*

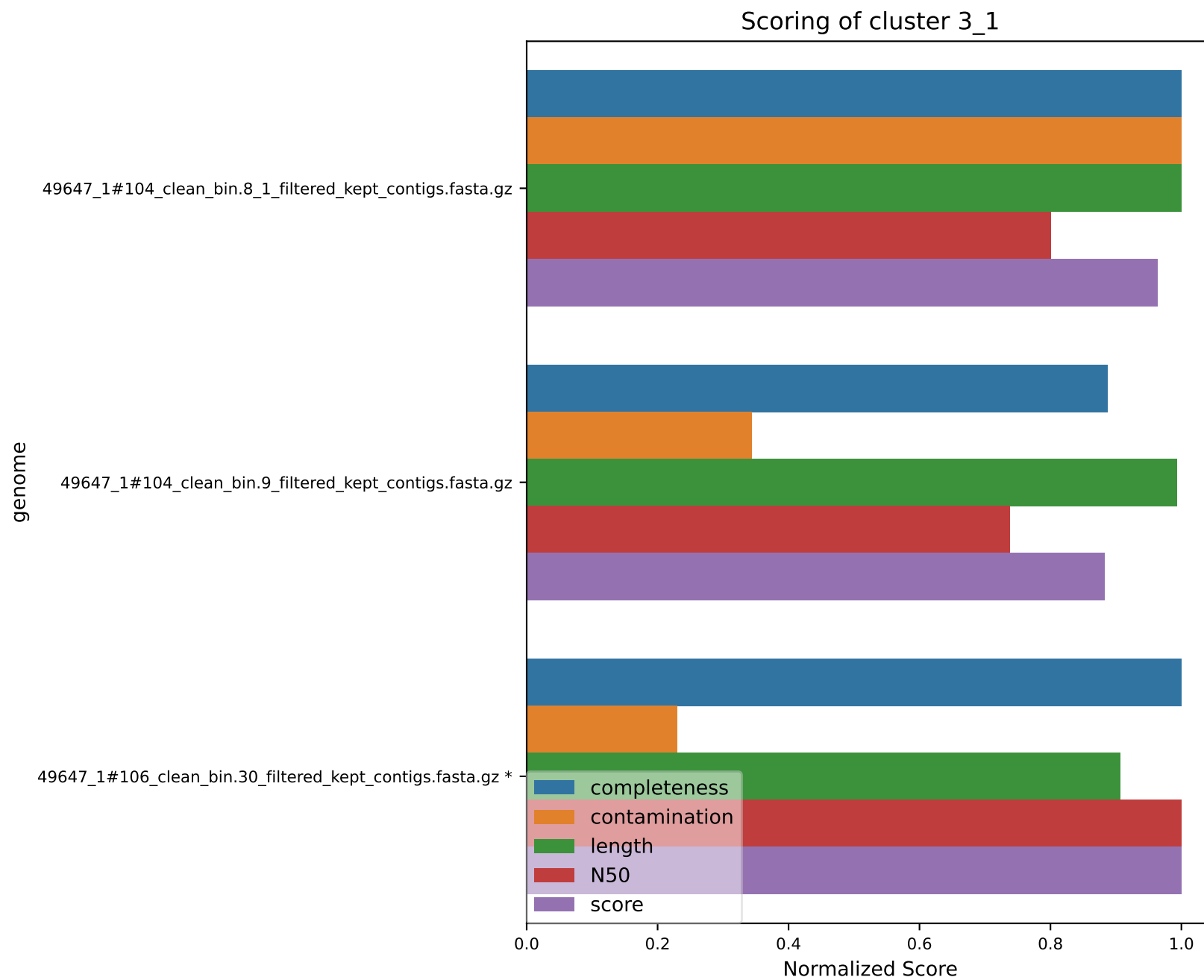


Scoring of cluster 2\_0

genome

49647\_1#2\_clean\_bin.19.orig\_filtered\_kept\_contigs.fasta.gz \*

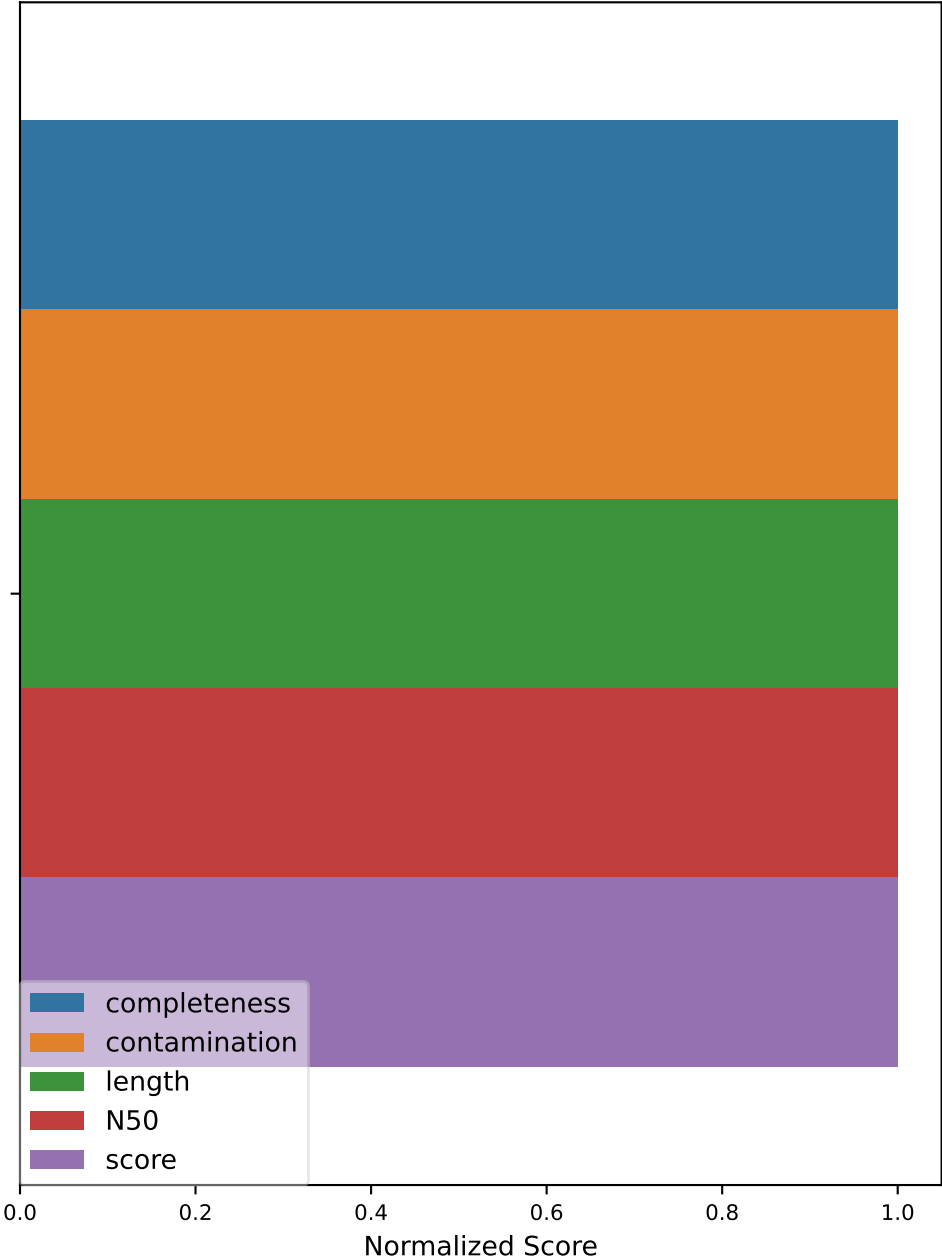




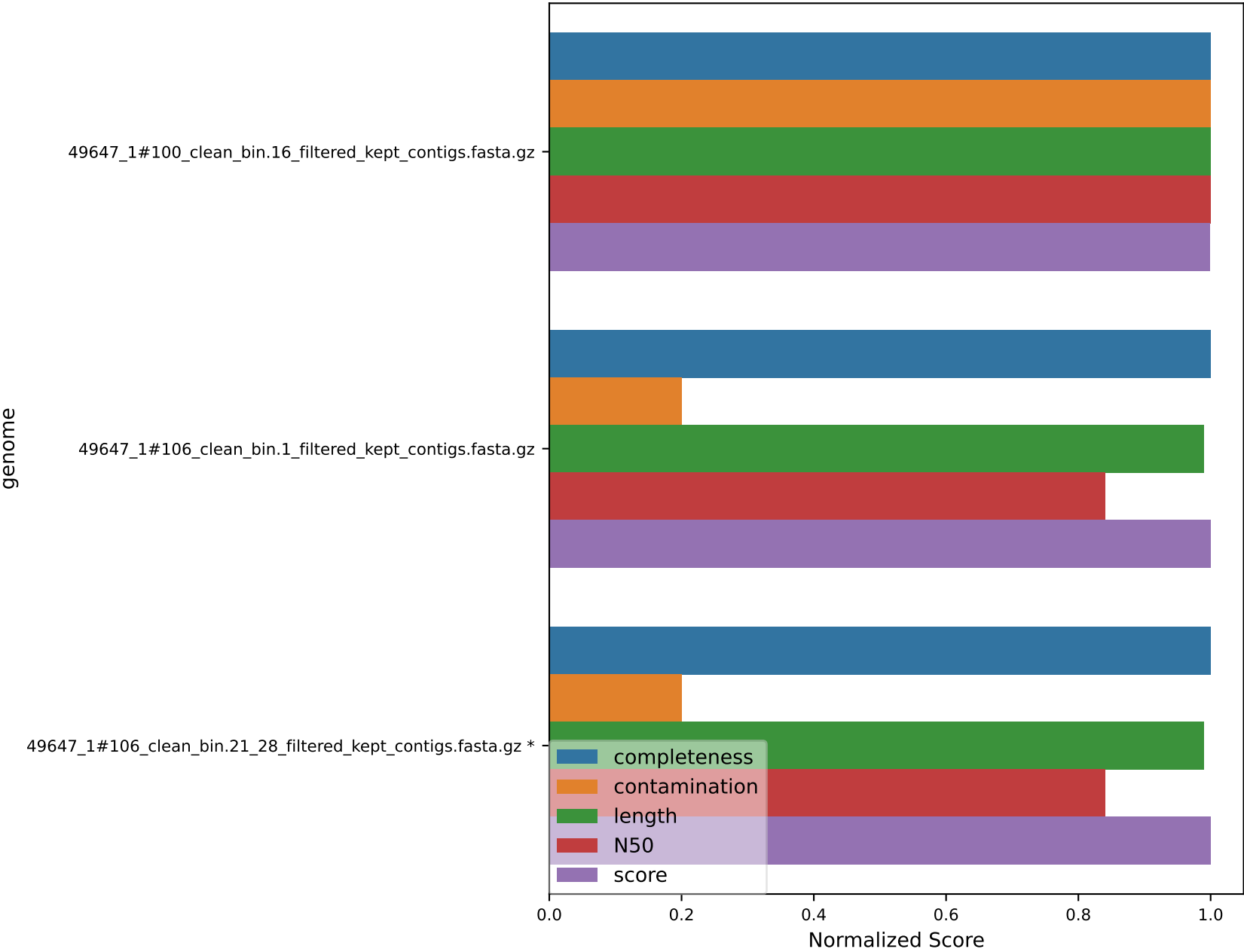
Scoring of cluster 4\_0

genome

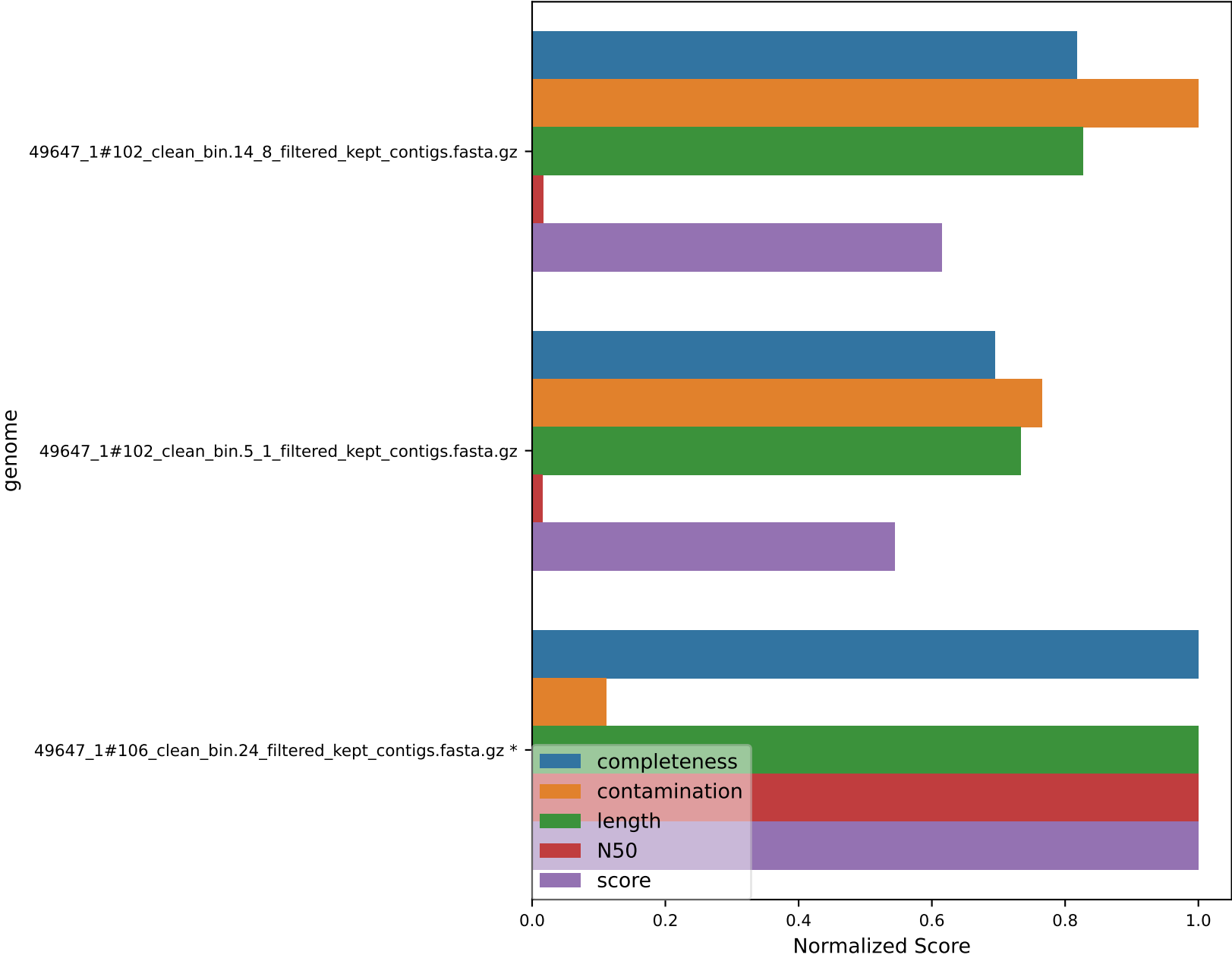
49647\_1#100\_clean\_bin.9\_filtered\_kept\_contigs.fasta.gz \*



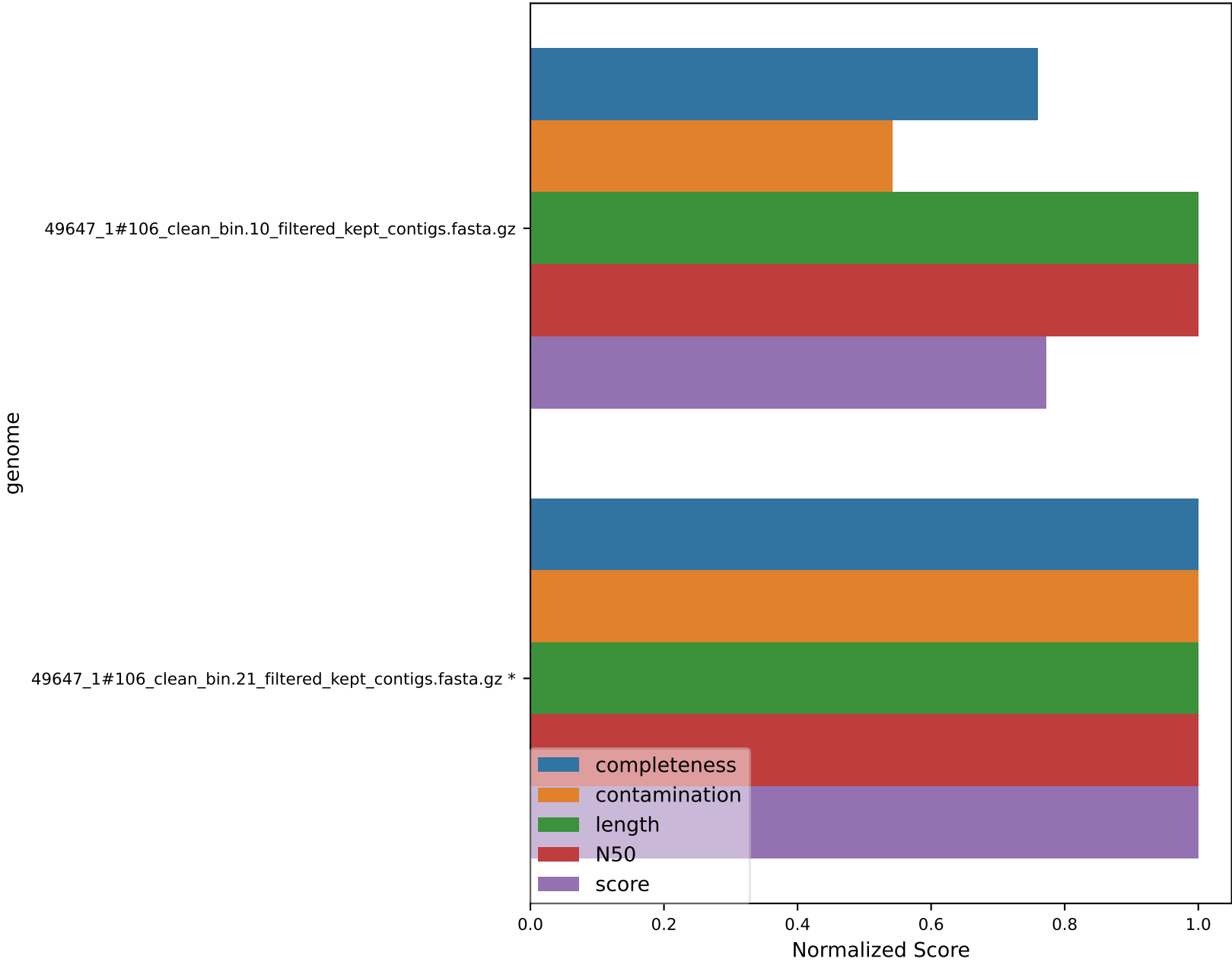
Scoring of cluster 5\_1



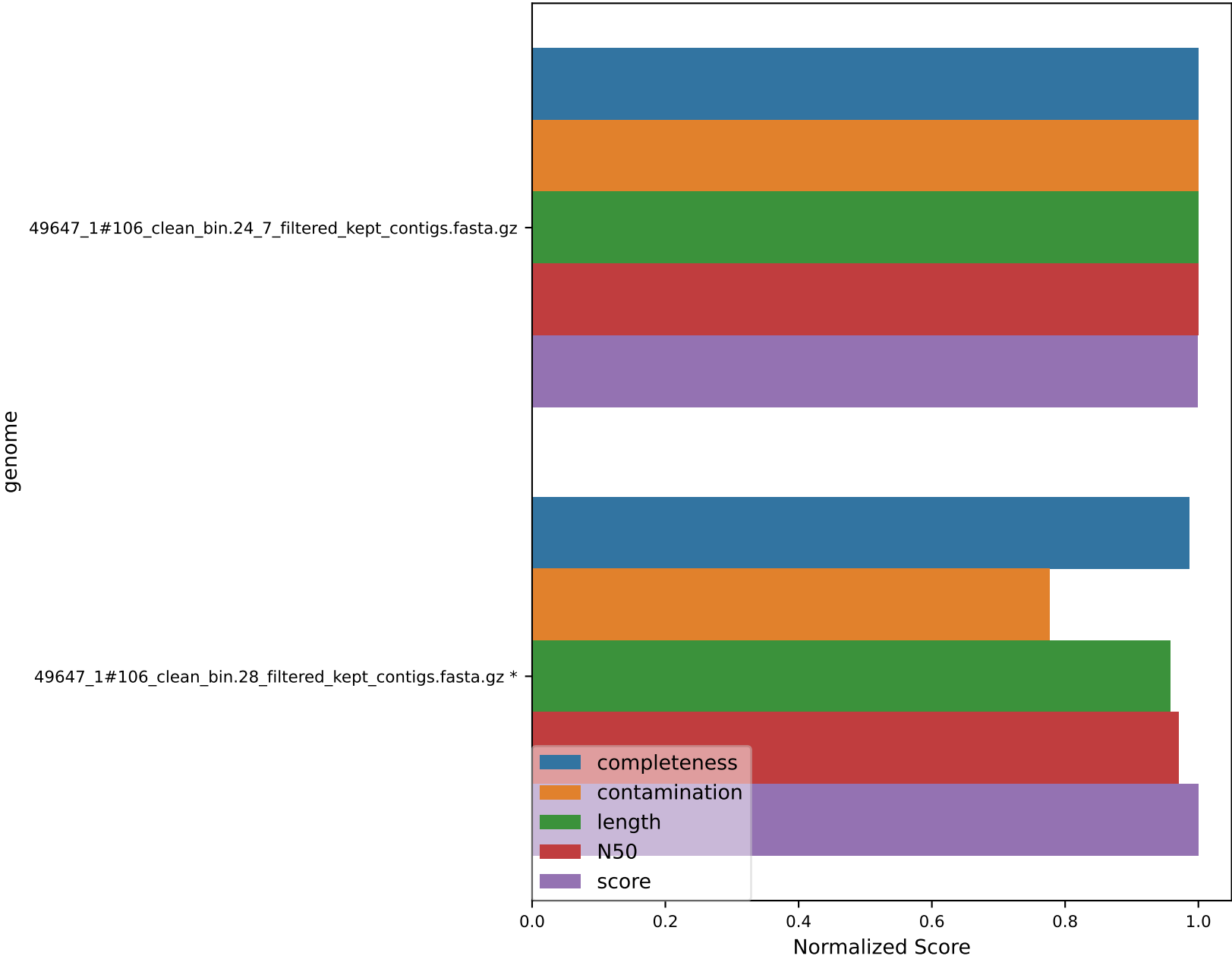
Scoring of cluster 6\_1



Scoring of cluster 7\_1

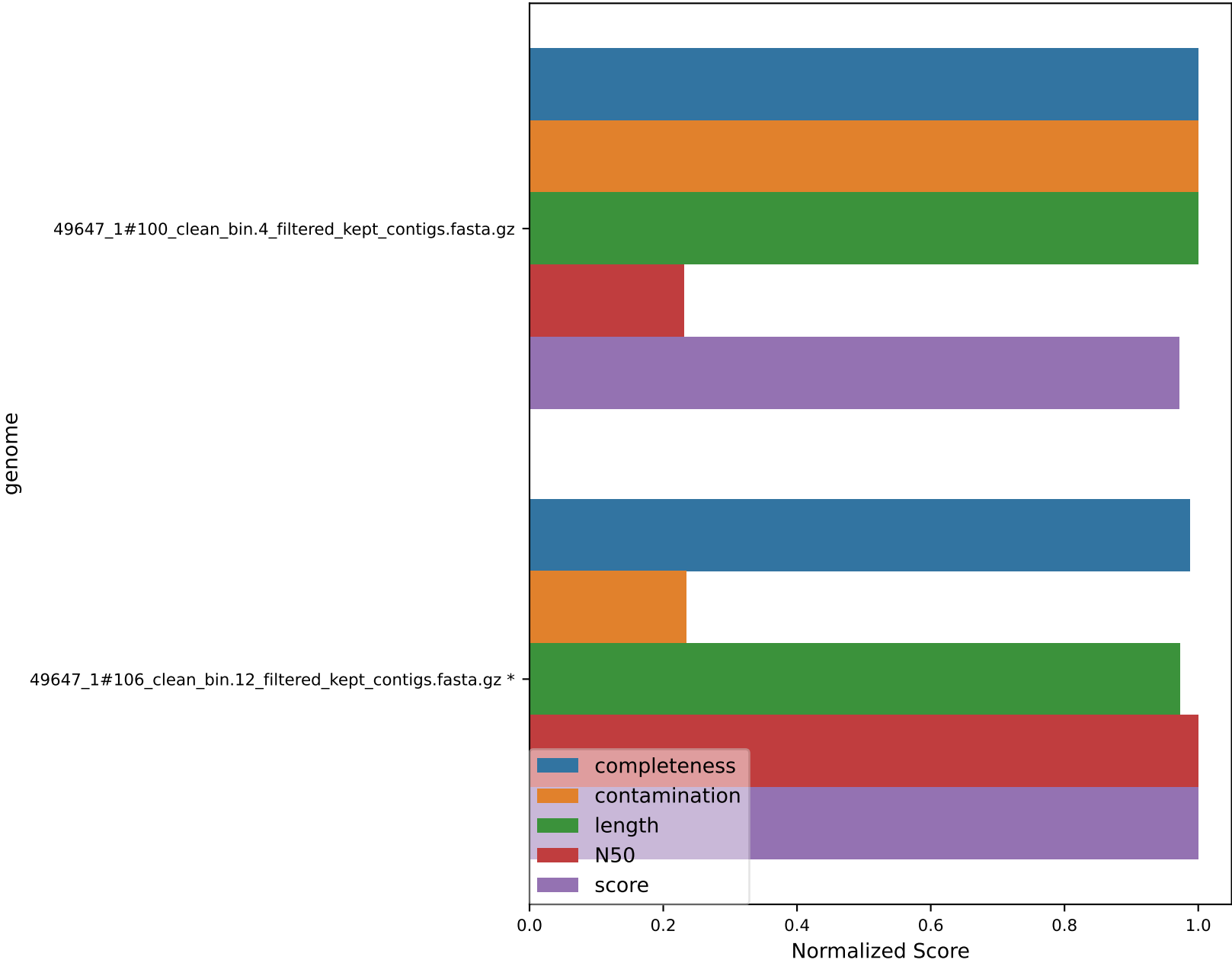


Scoring of cluster 8\_1





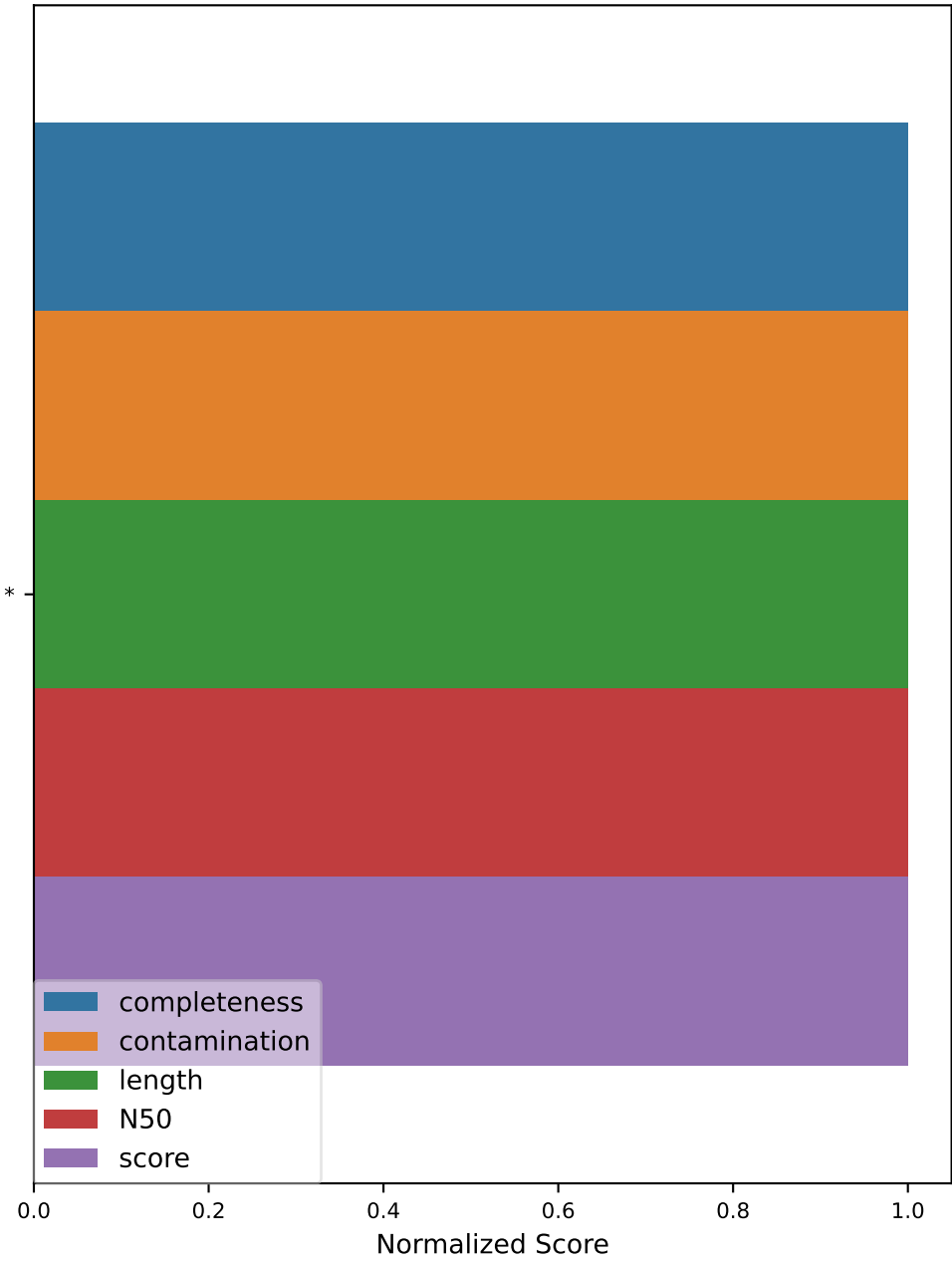
Scoring of cluster 9\_1



Scoring of cluster 10\_0

genome

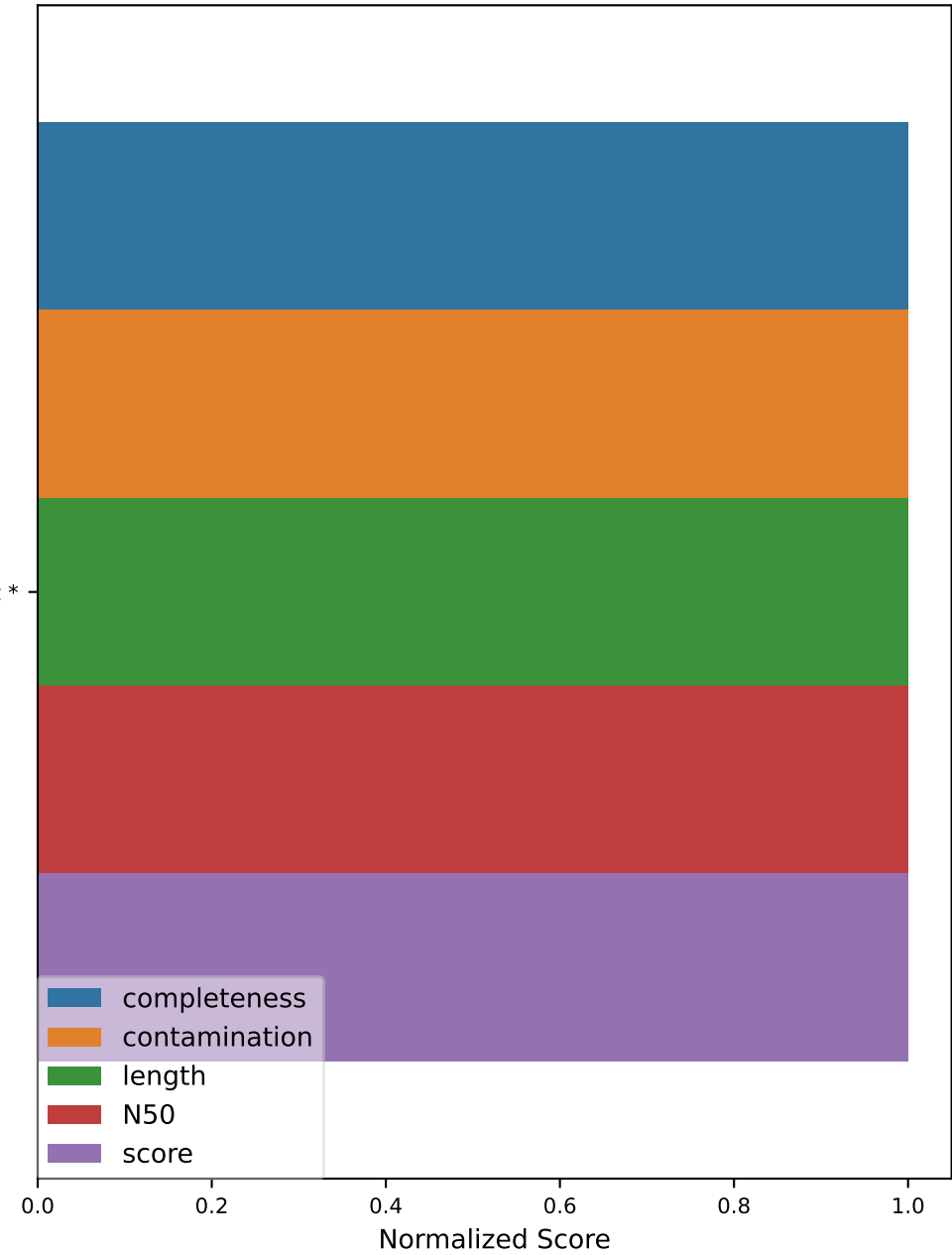
49647\_1#100\_clean\_bin.18\_filtered\_kept\_contigs.fasta.gz \*



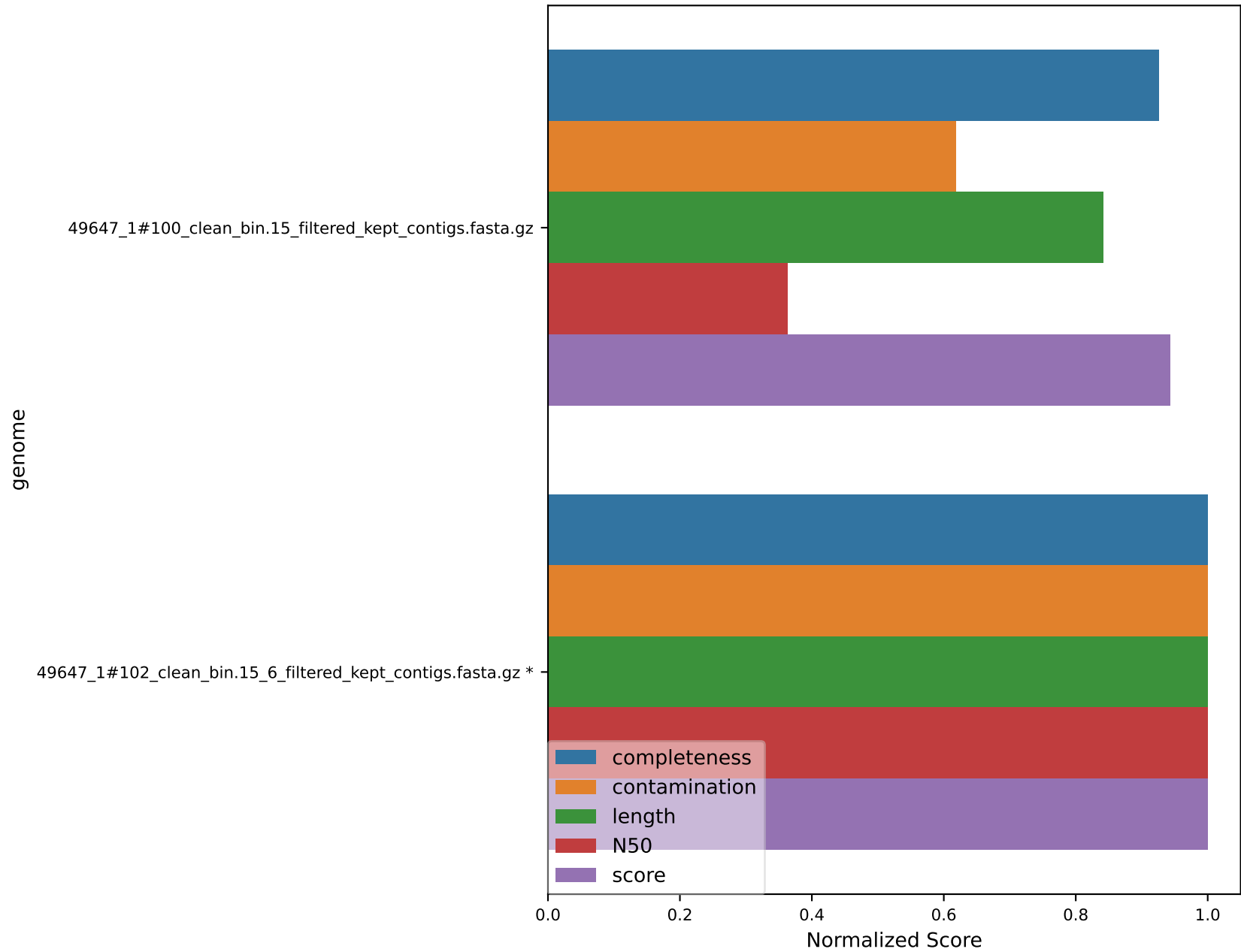
Scoring of cluster 11\_0

genome

49647\_1#106\_clean\_bin.26\_24\_filtered\_kept\_contigs.fasta.gz \*



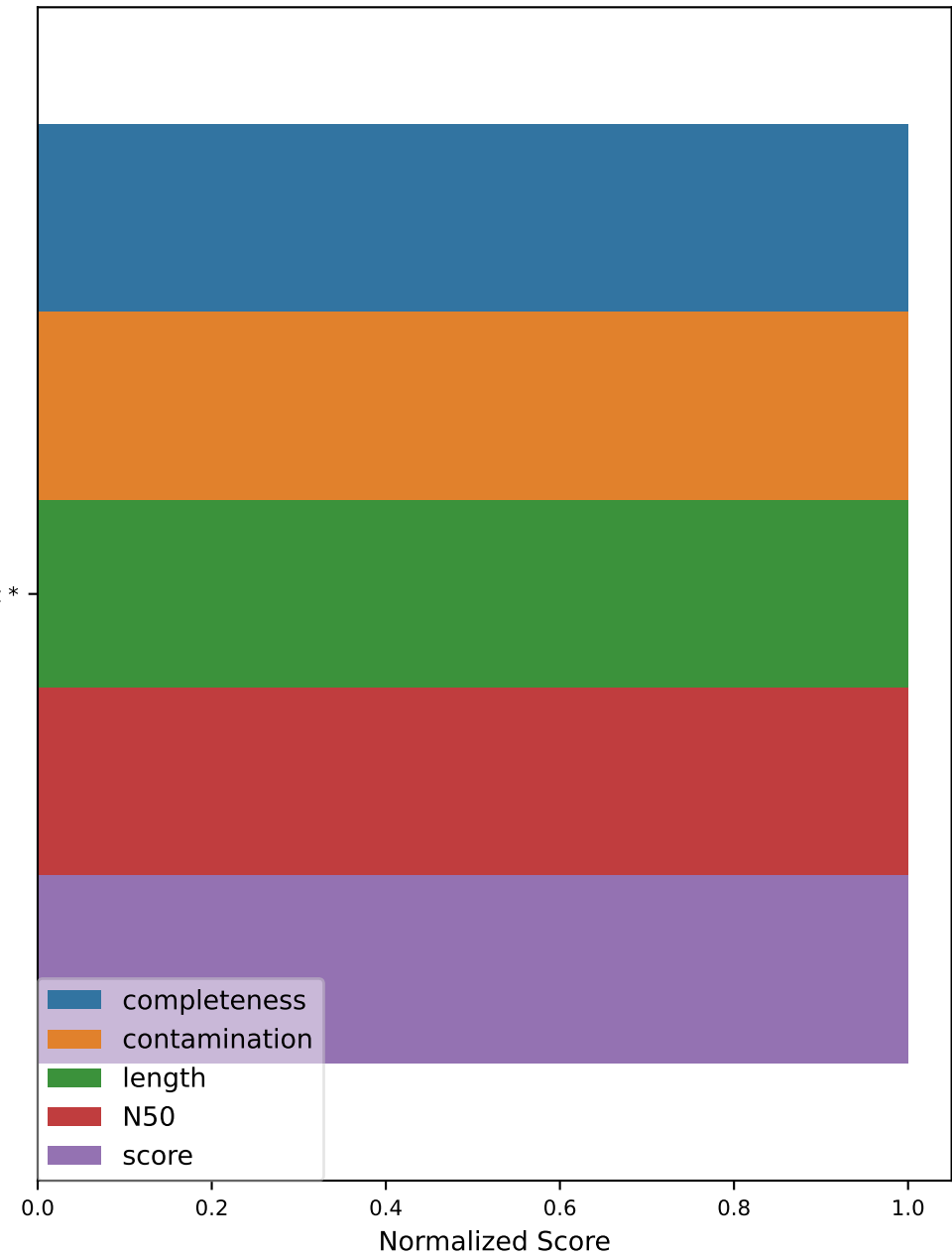
Scoring of cluster 12\_1



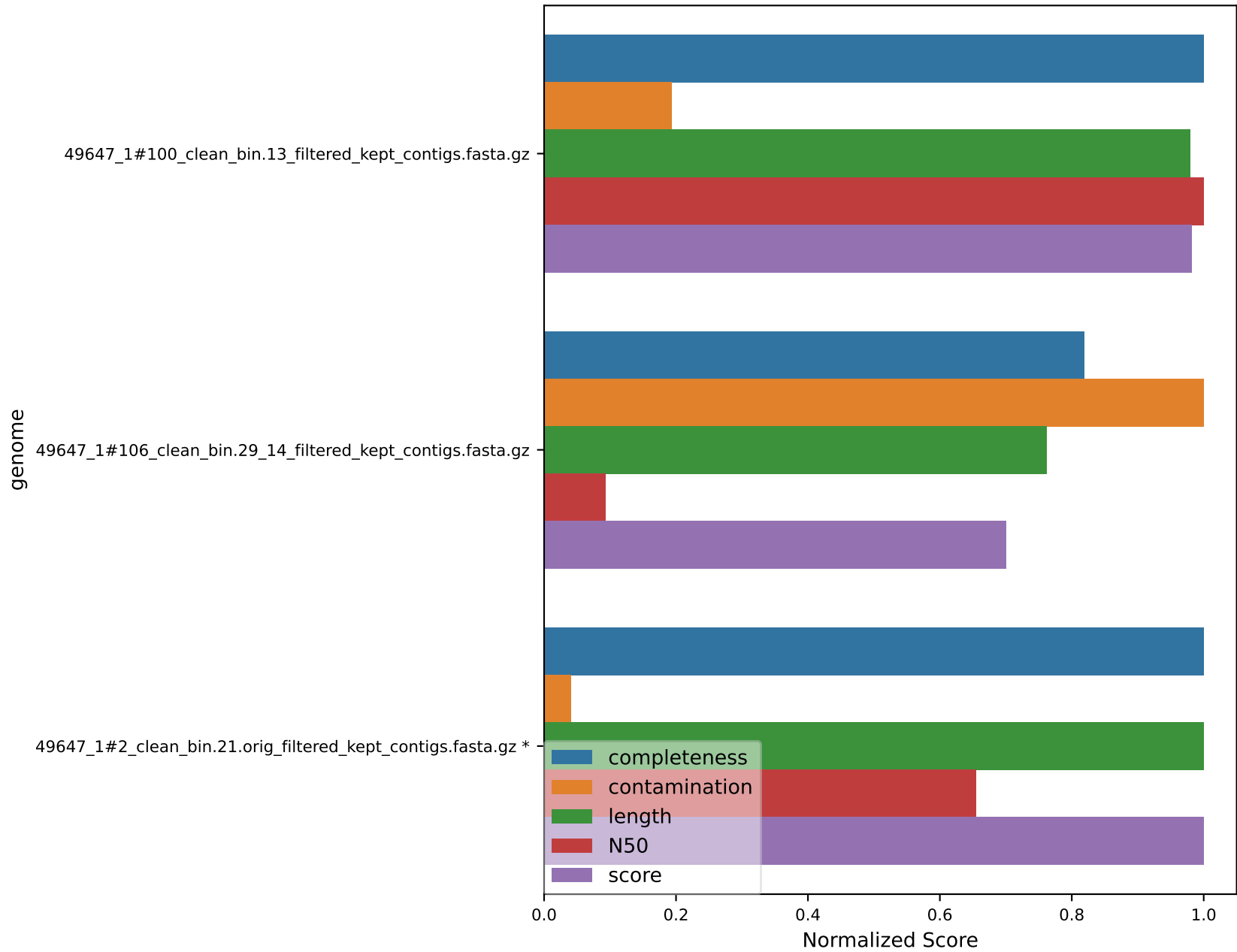
Scoring of cluster 13\_0

genome

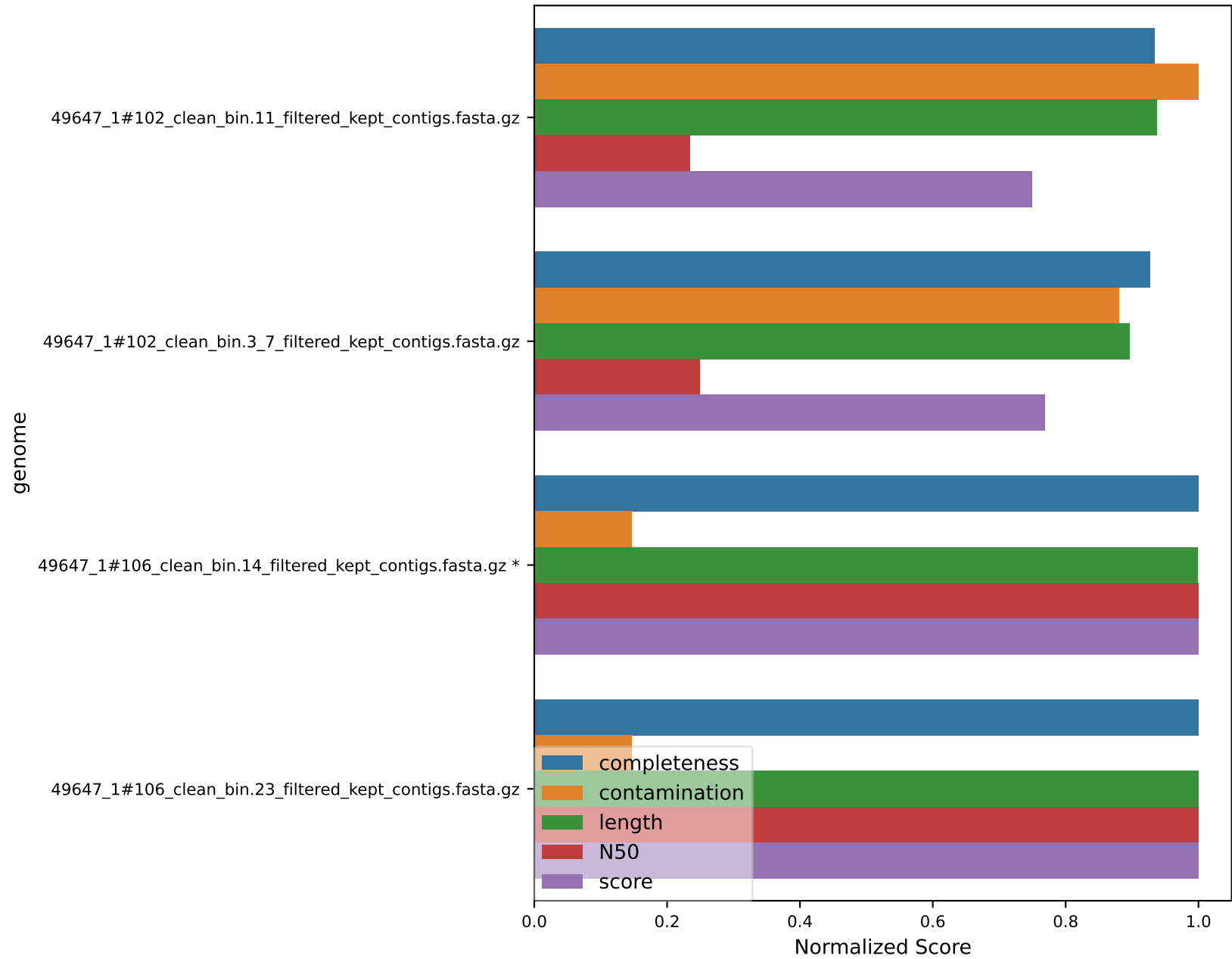
49647\_1#2\_clean\_bin.20.orig\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 14\_1



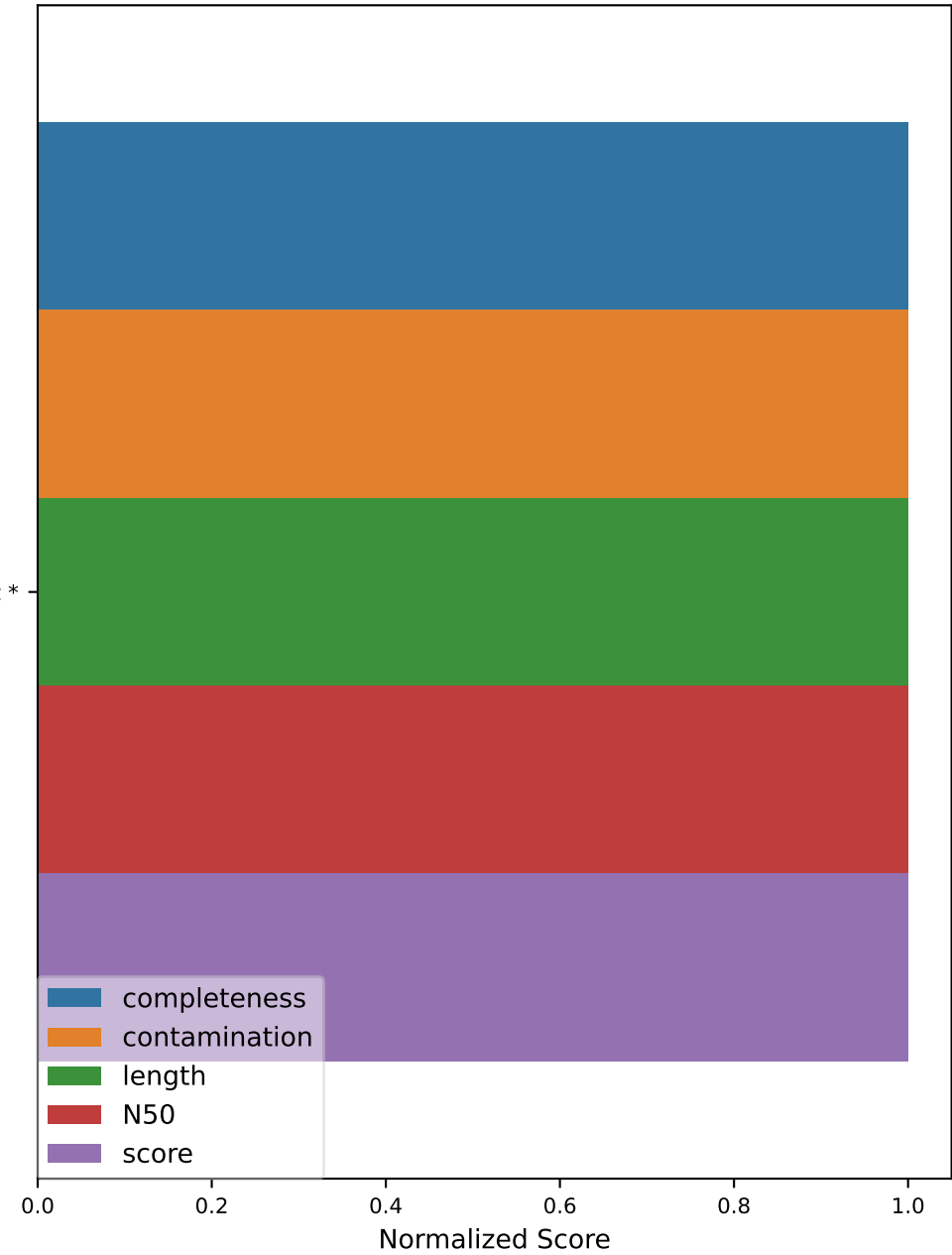
# Scoring of cluster 15\_1



Scoring of cluster 16\_0

genome

49647\_1#2\_clean\_bin.15.strict\_filtered\_kept\_contigs.fasta.gz \*

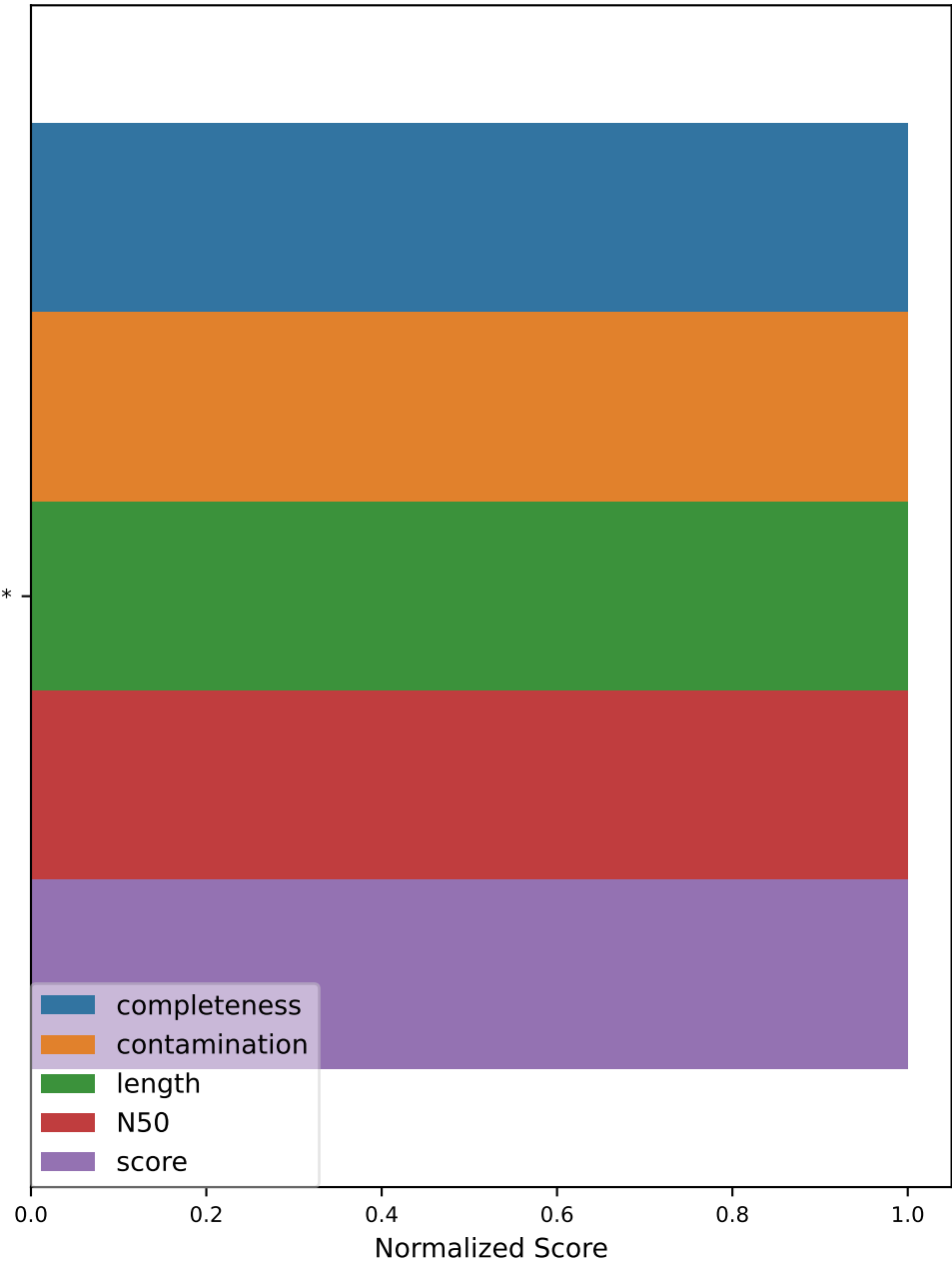




Scoring of cluster 17\_0

genome

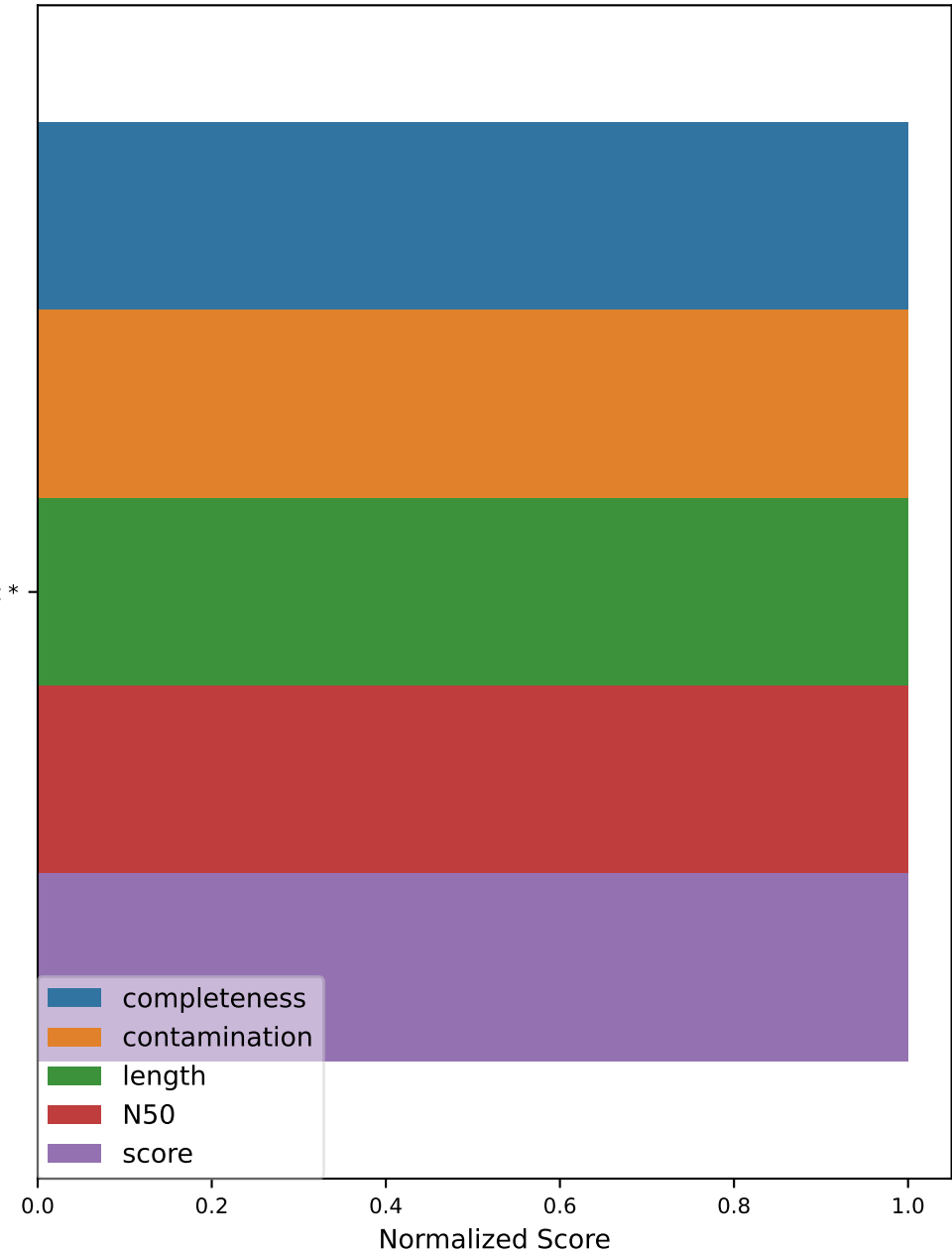
49647\_1#2\_clean\_bin.26.orig\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 18\_0

genome

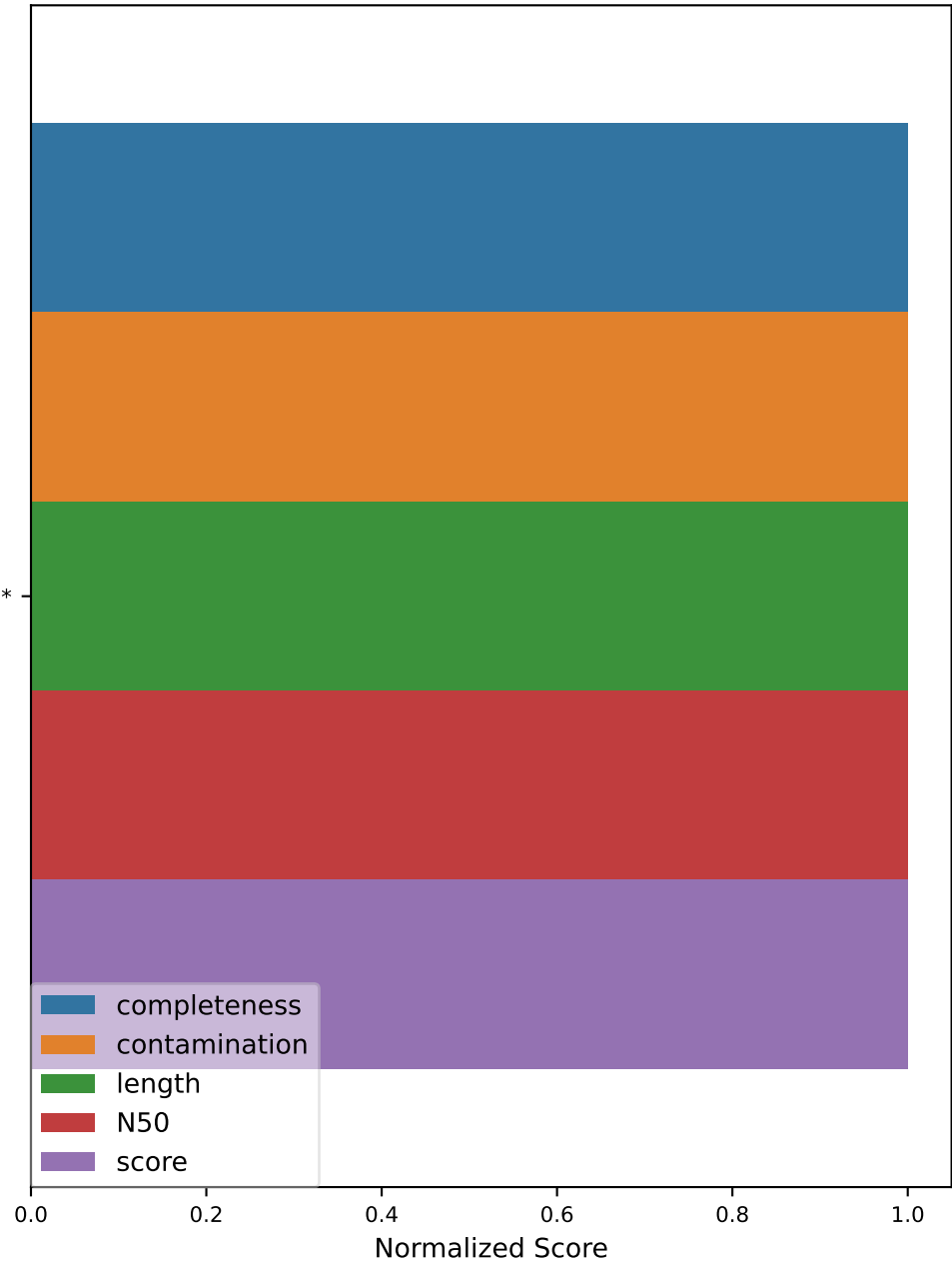
49647\_1#2\_clean\_bin.8.permissive\_filtered\_kept\_contigs.fasta.gz \*



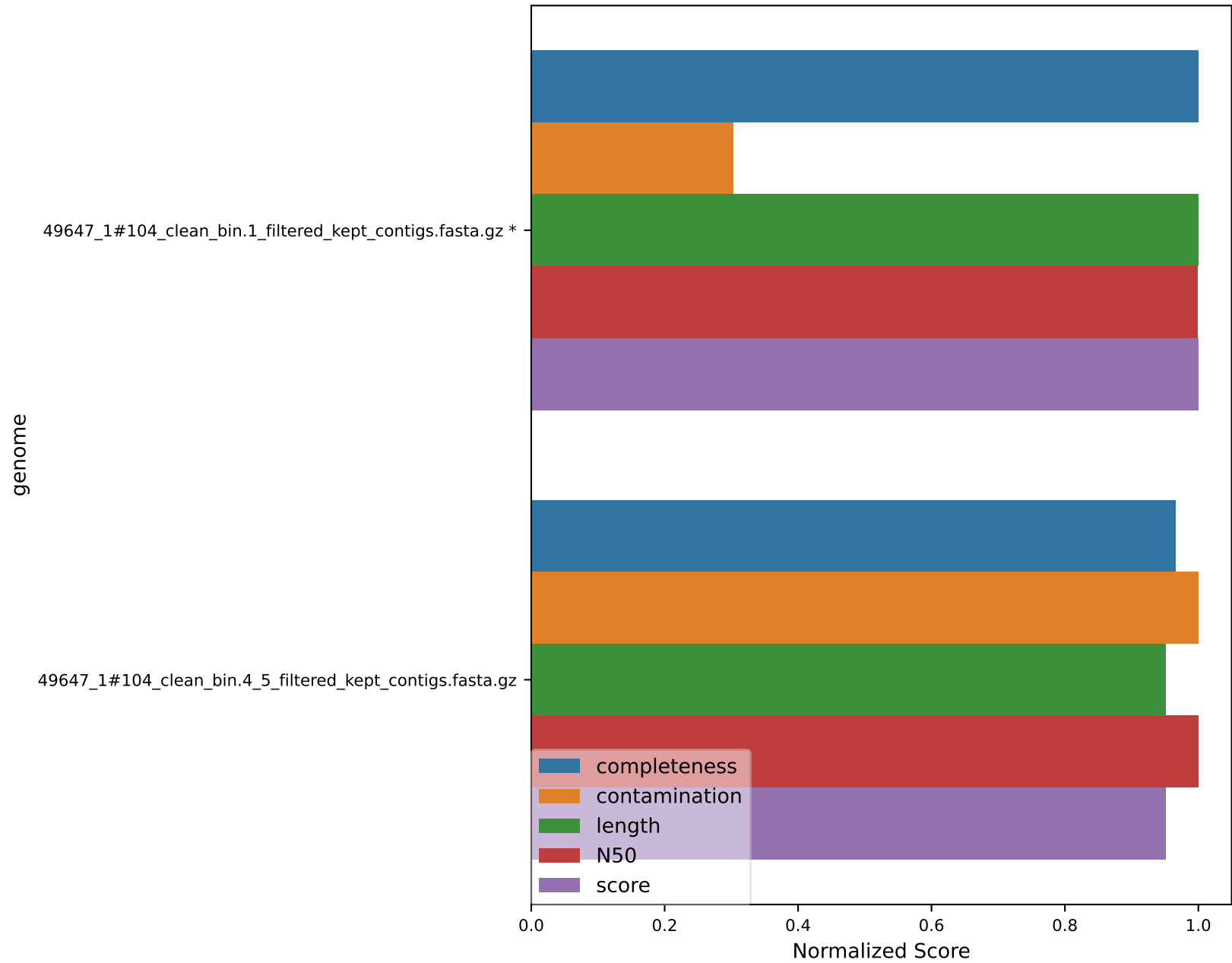
Scoring of cluster 19\_0

genome

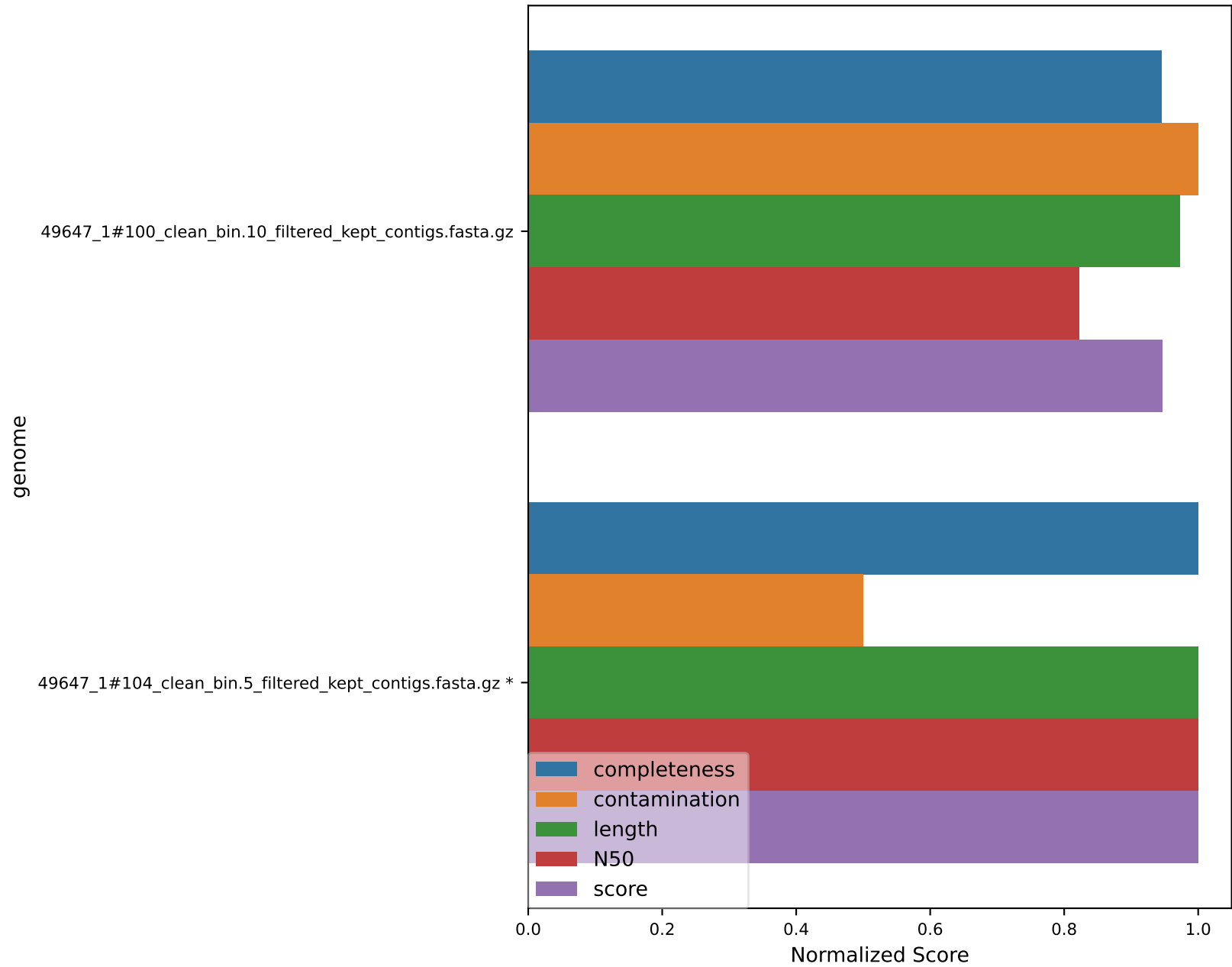
49647\_1#100\_clean\_bin.1\_filtered\_kept\_contigs.fasta.gz \*



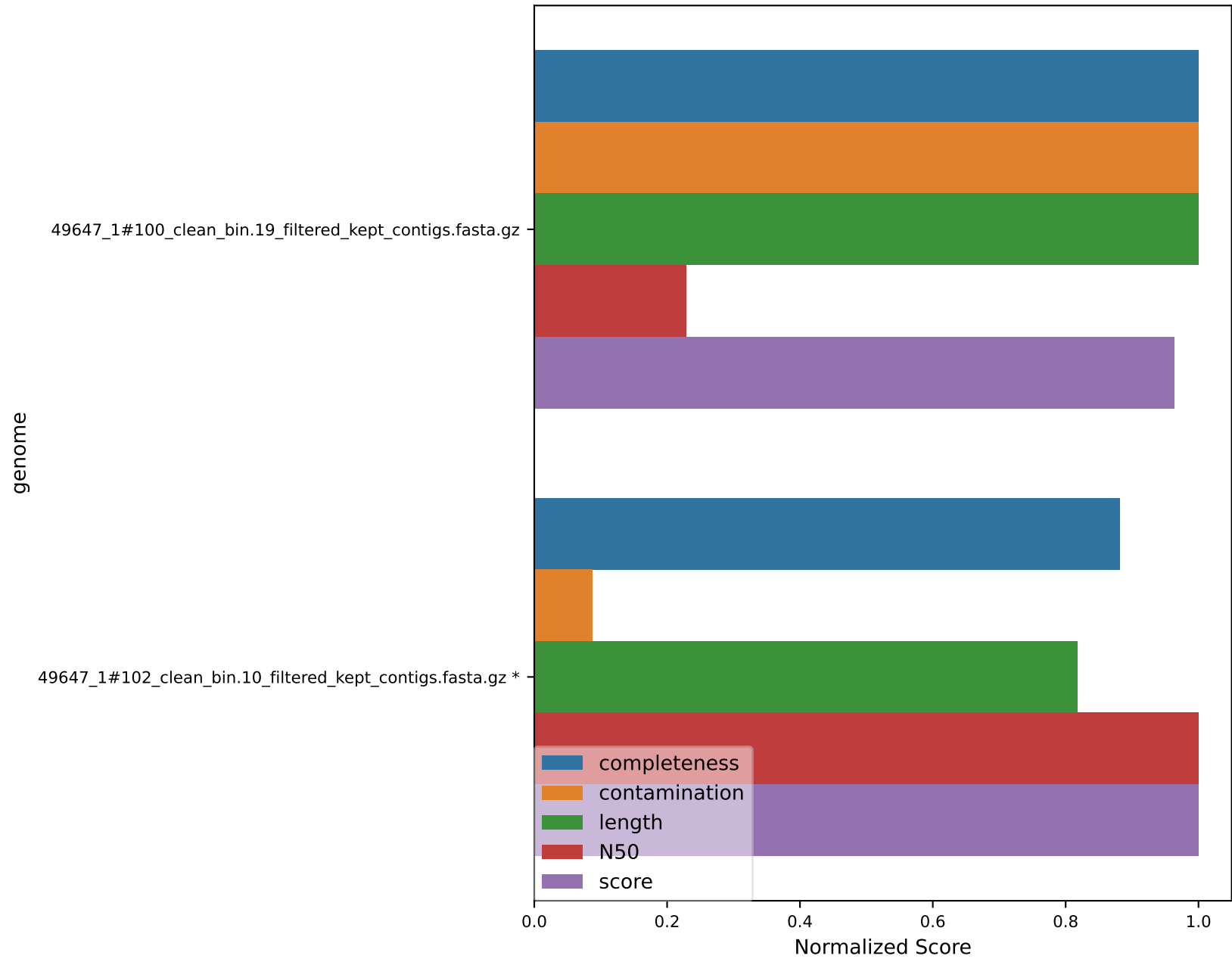
Scoring of cluster 20\_1



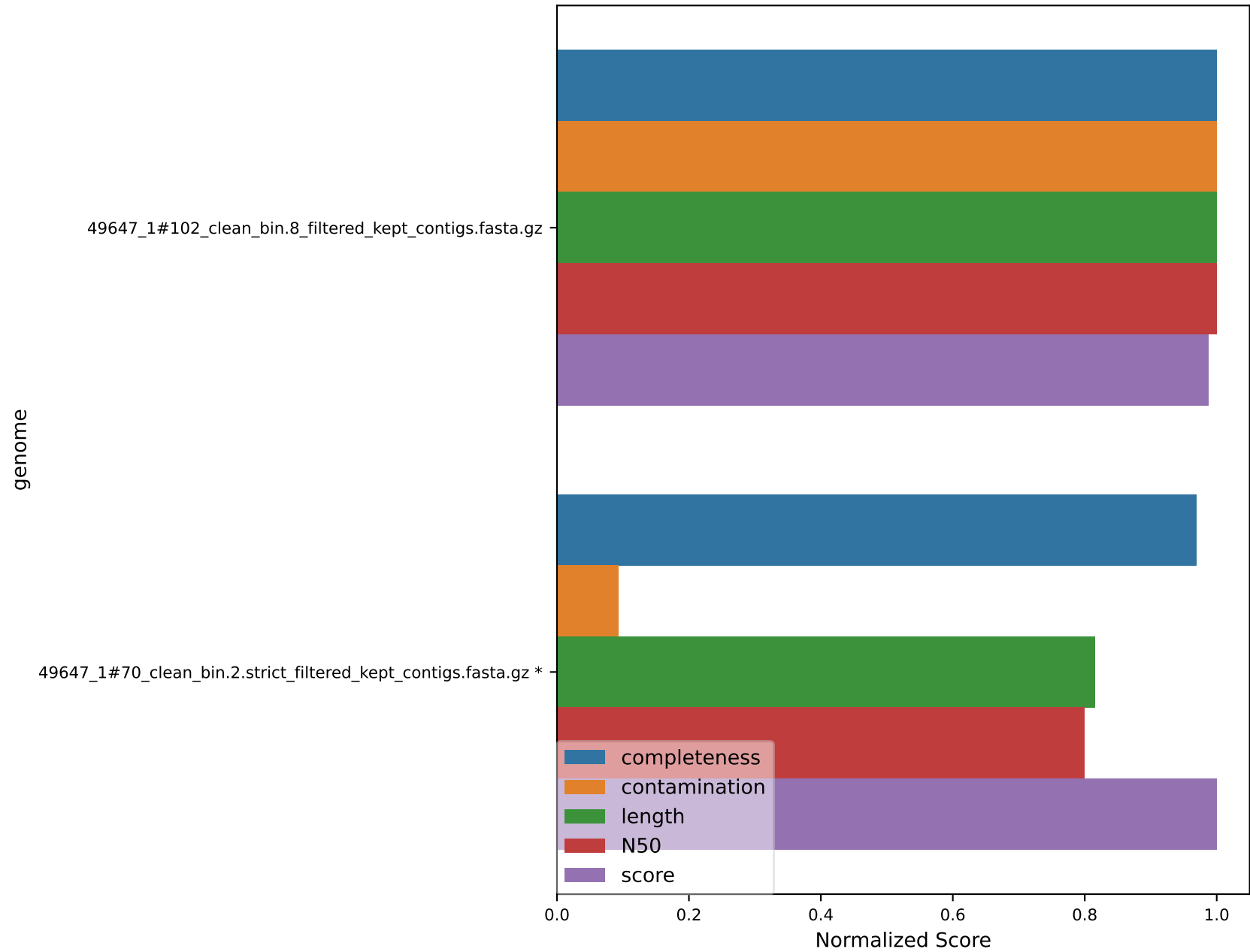
Scoring of cluster 21\_1



Scoring of cluster 22\_1



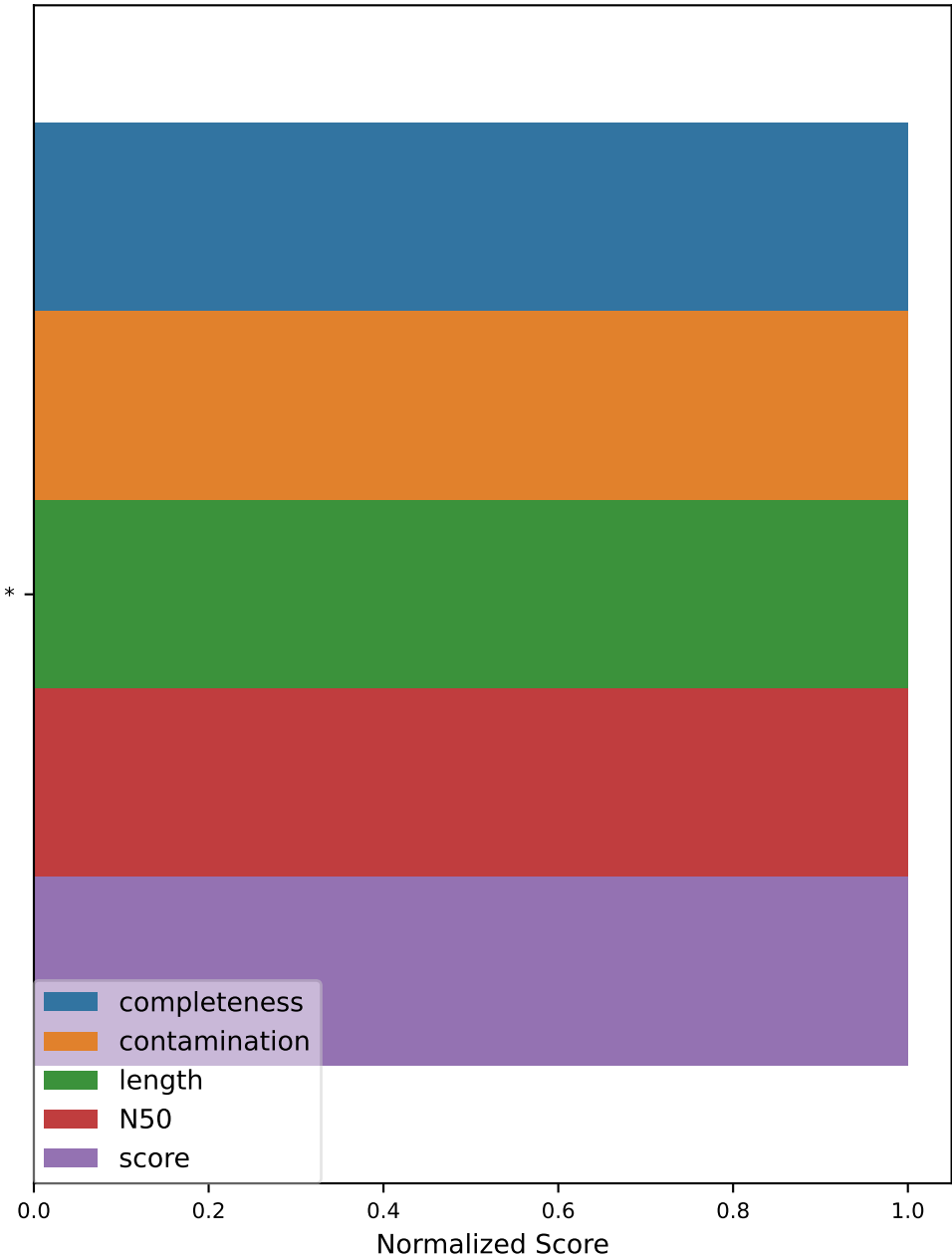
Scoring of cluster 23\_1



Scoring of cluster 23\_2

genome

49647\_1#2\_clean\_bin.30.orig\_filtered\_kept\_contigs.fasta.gz \*

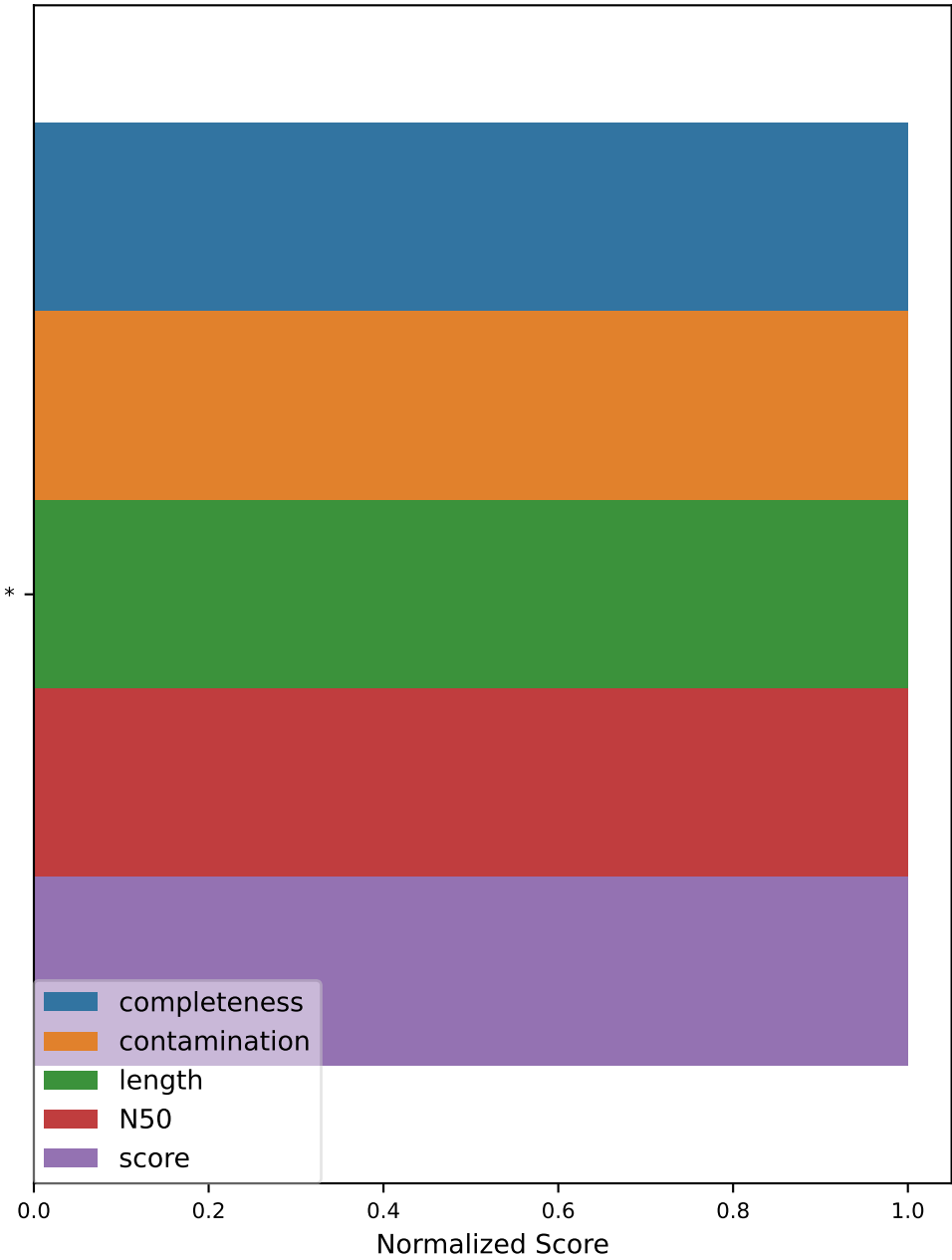




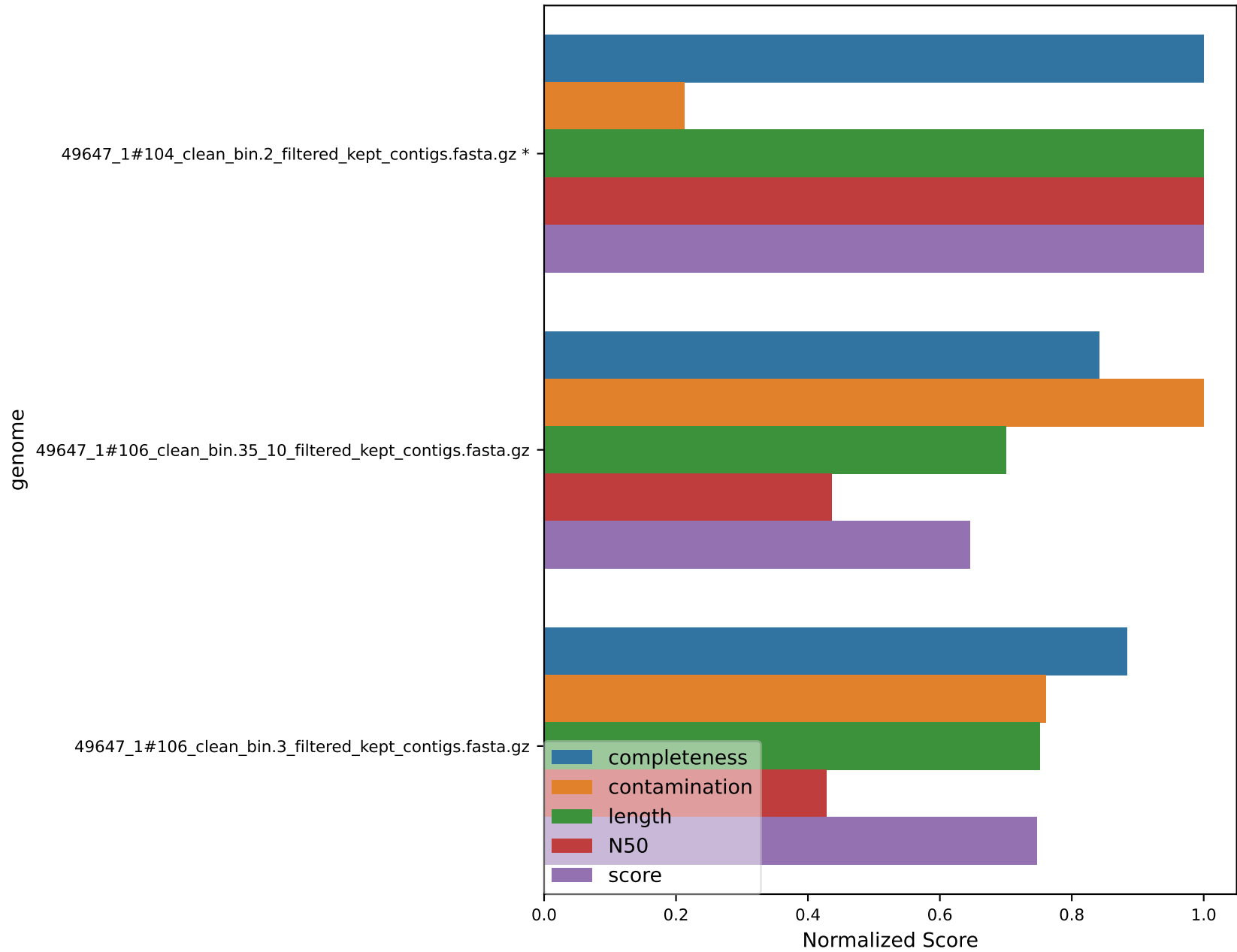
Scoring of cluster 24\_0

genome

49647\_1#102\_clean\_bin.12\_filtered\_kept\_contigs.fasta.gz \*



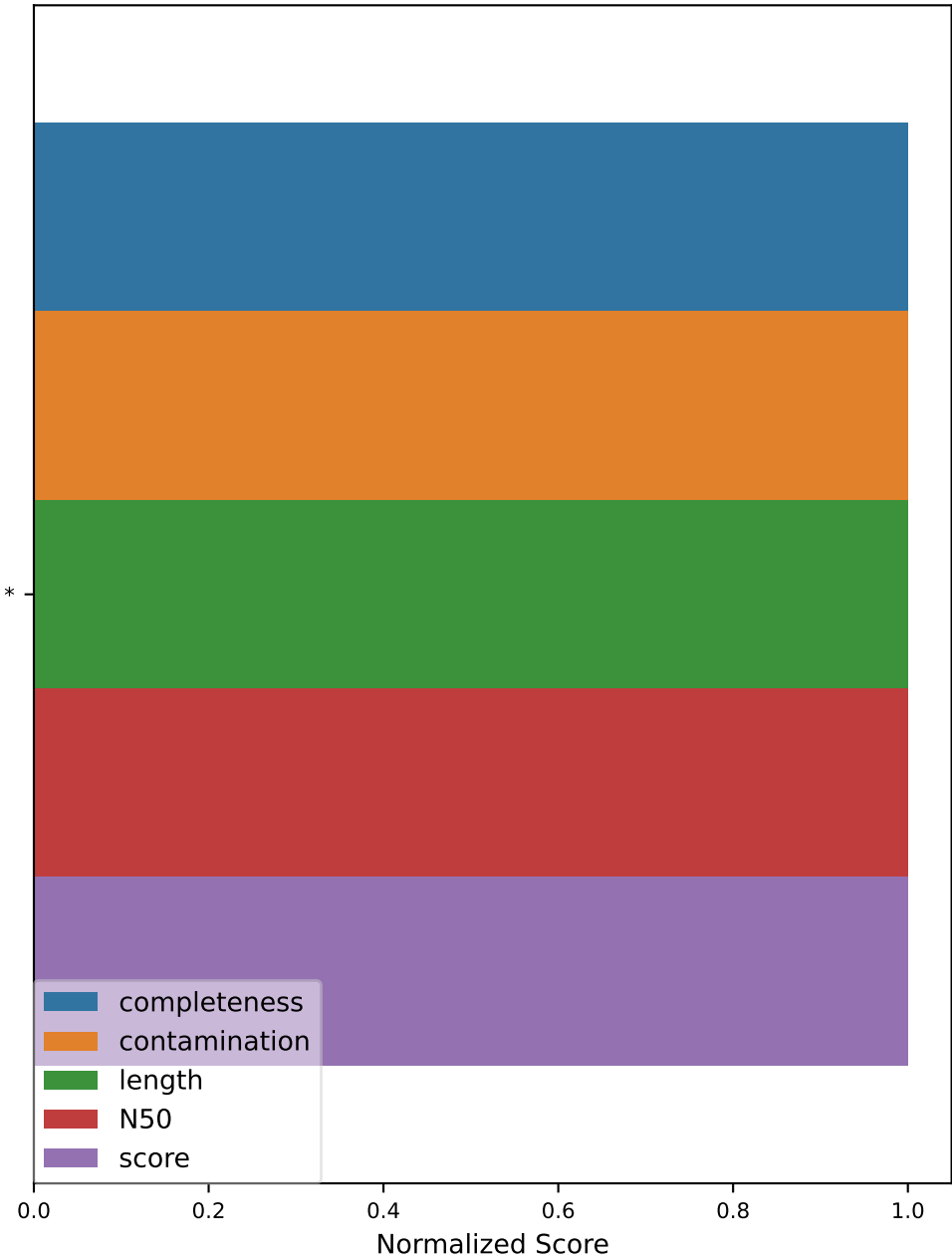
Scoring of cluster 25\_1



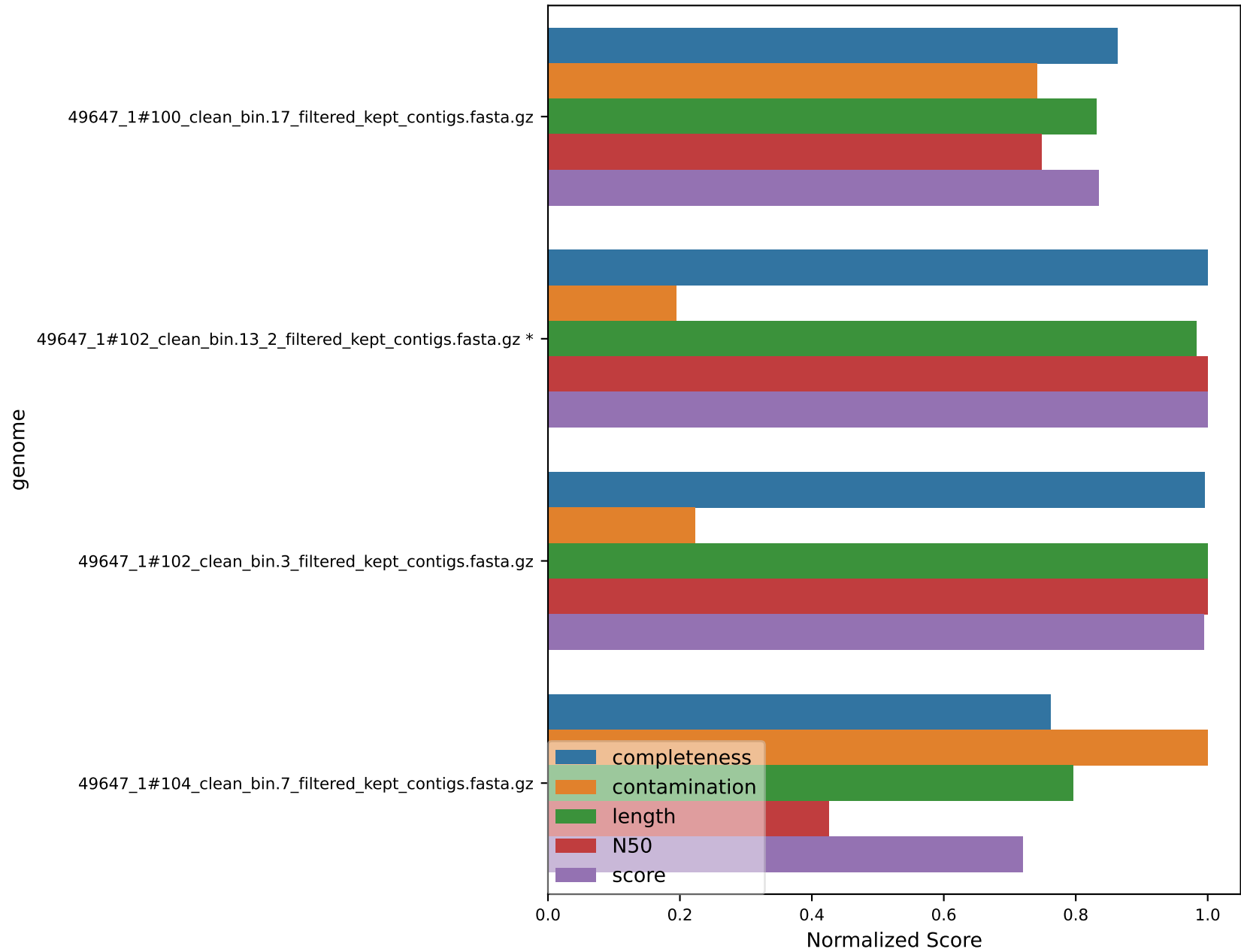
Scoring of cluster 26\_0

genome

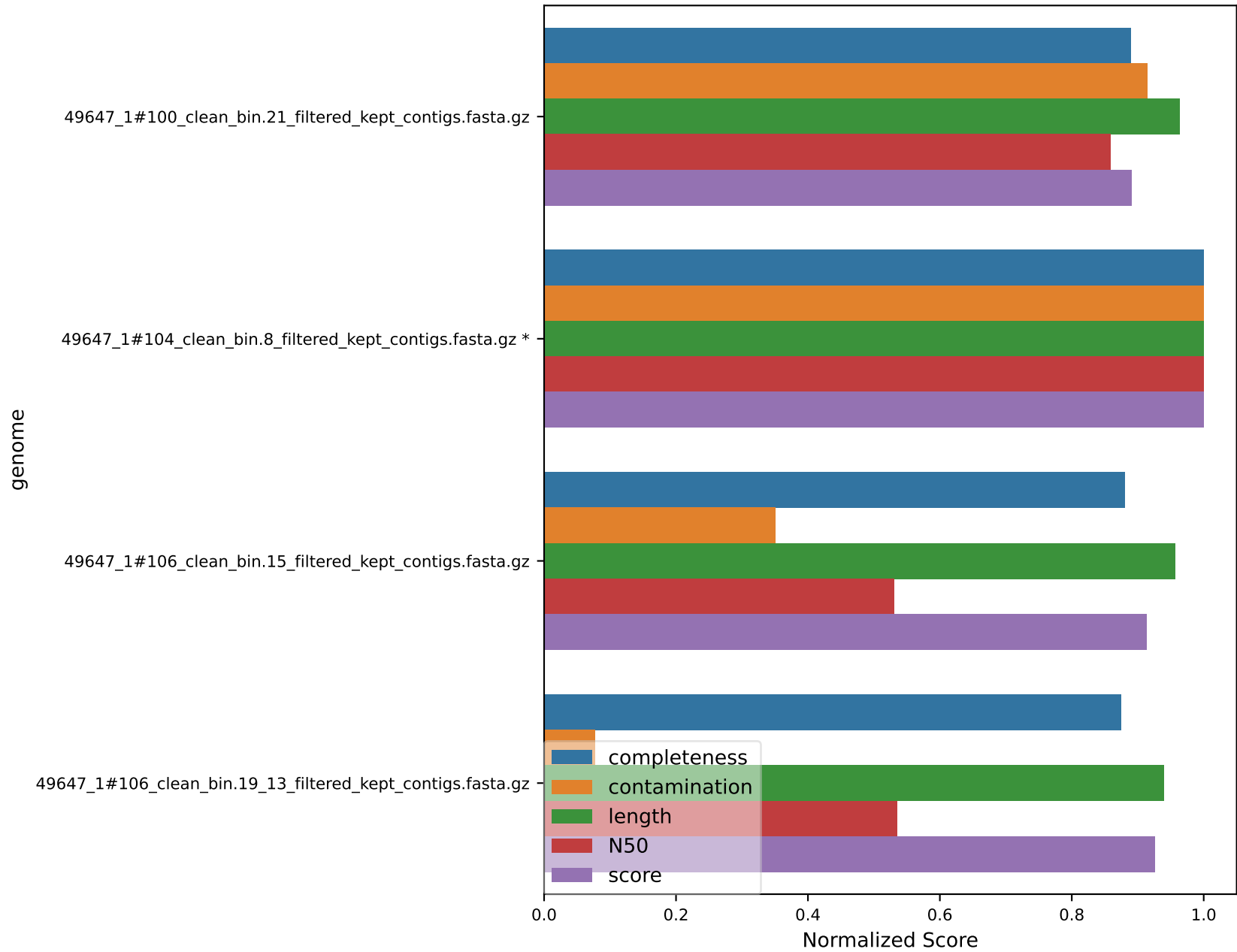
49647\_1#100\_clean\_bin.22\_filtered\_kept\_contigs.fasta.gz \*



# Scoring of cluster 27\_1



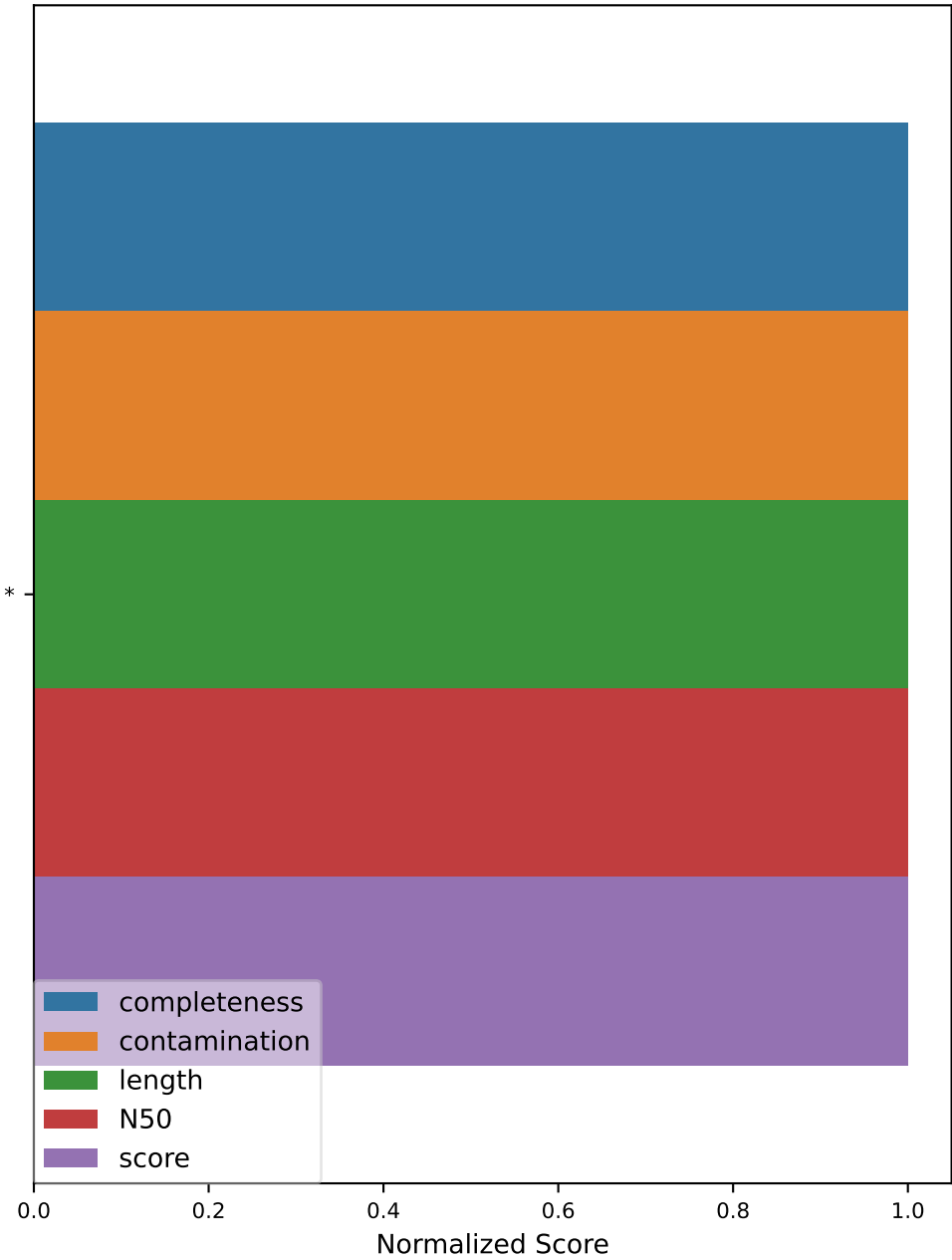
# Scoring of cluster 28\_1



Scoring of cluster 29\_0

genome

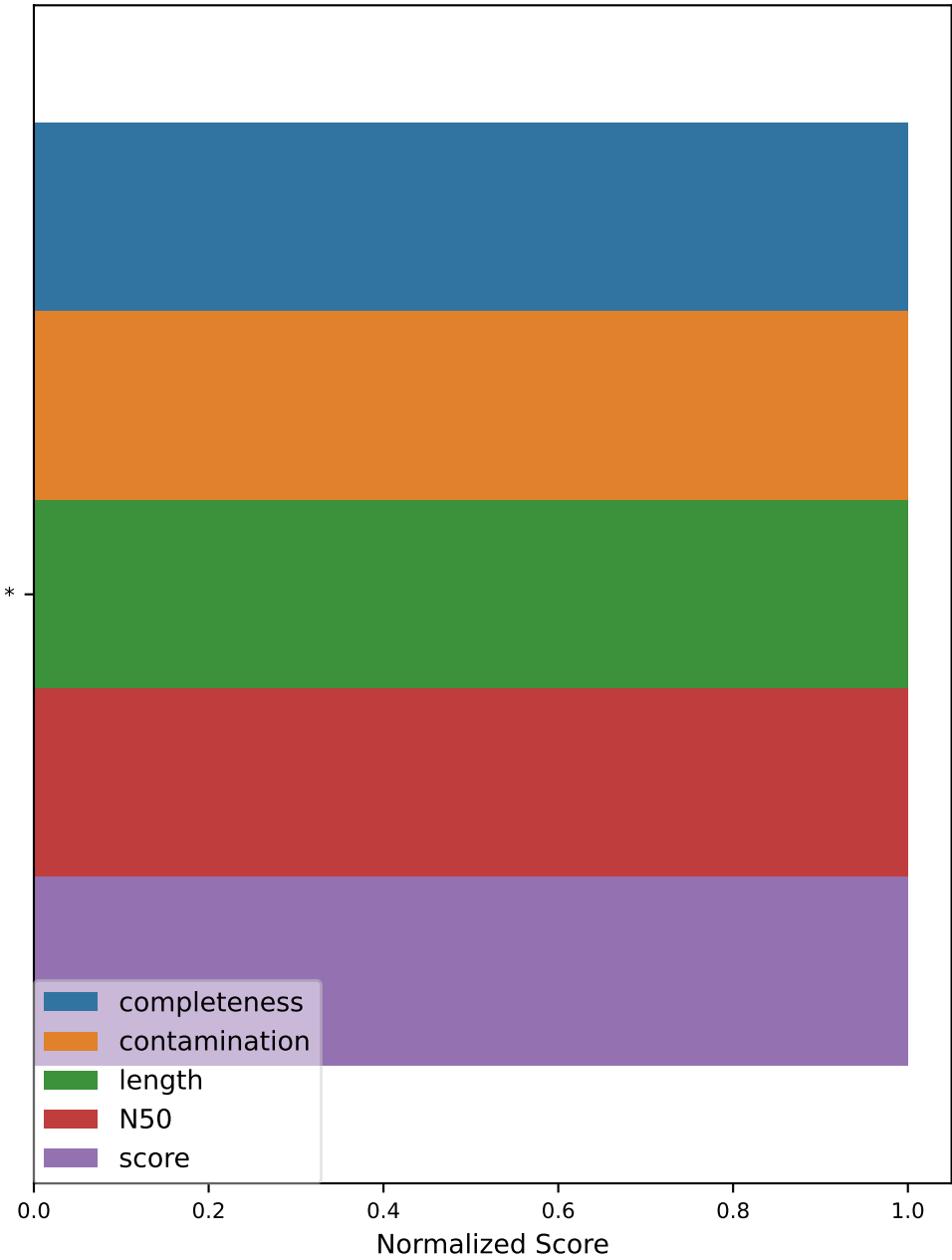
49647\_1#102\_clean\_bin.6\_3\_filtered\_kept\_contigs.fasta.gz \*



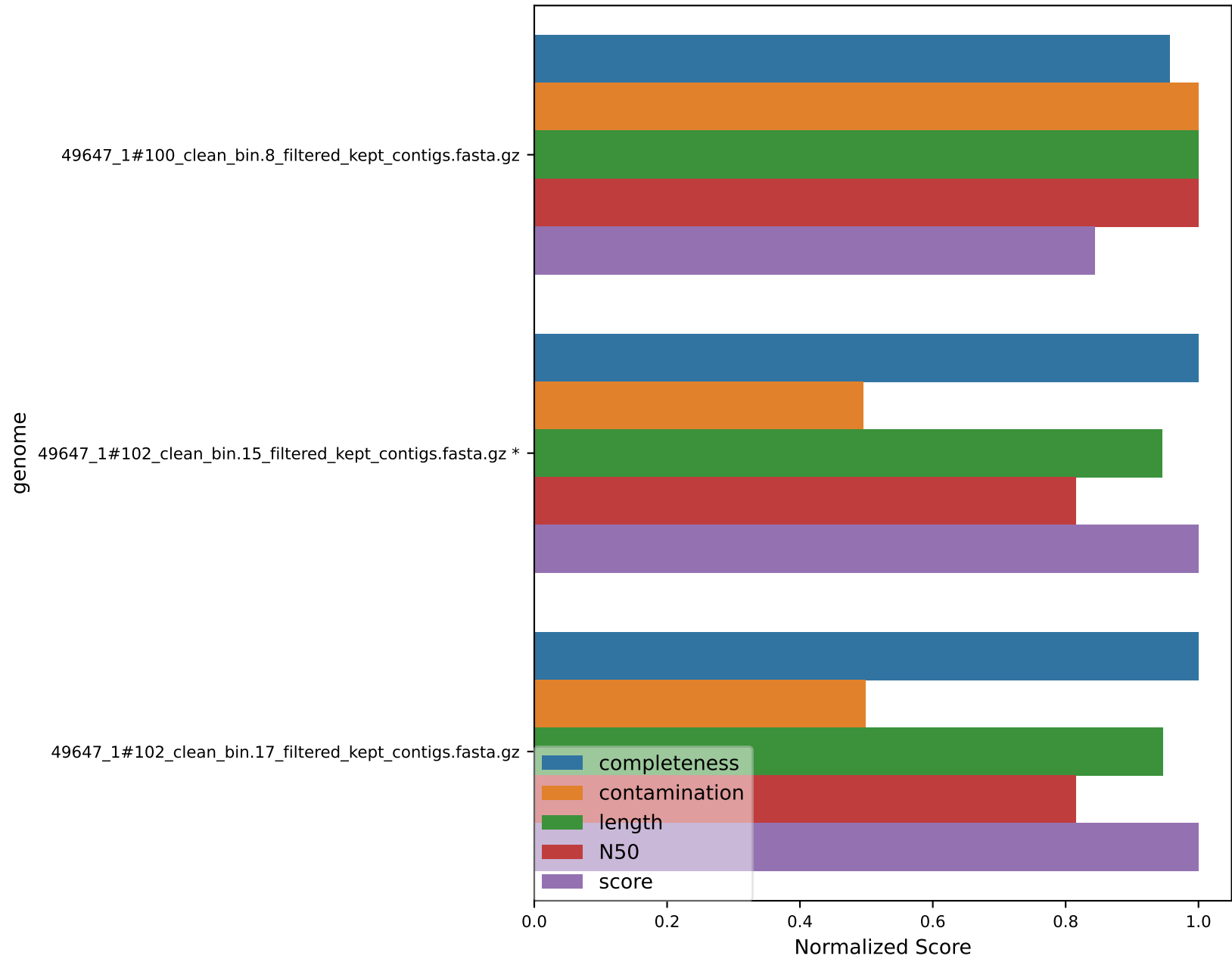
Scoring of cluster 30\_0

genome

49647\_1#100\_clean\_bin.20\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 31\_1

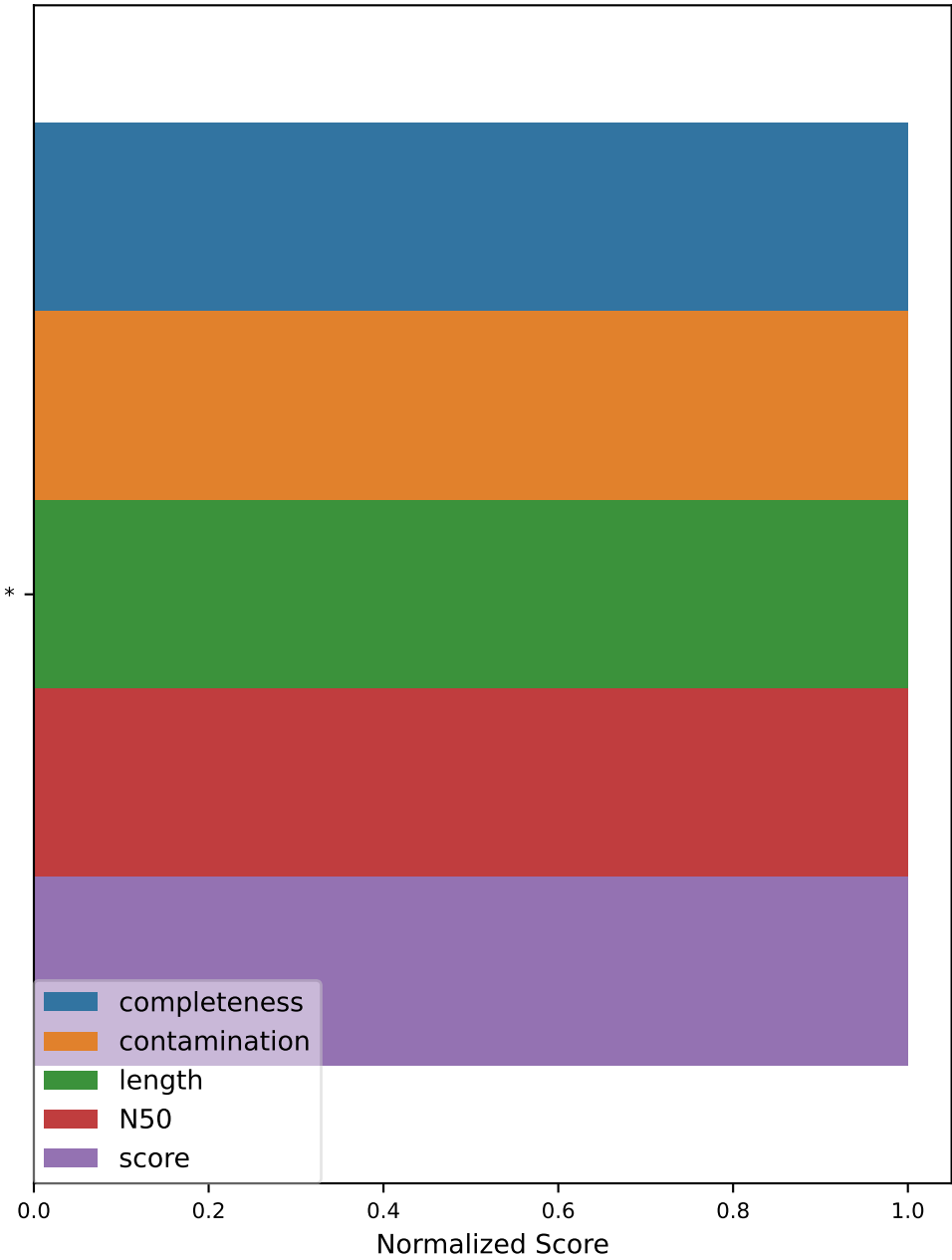




Scoring of cluster 32\_0

genome

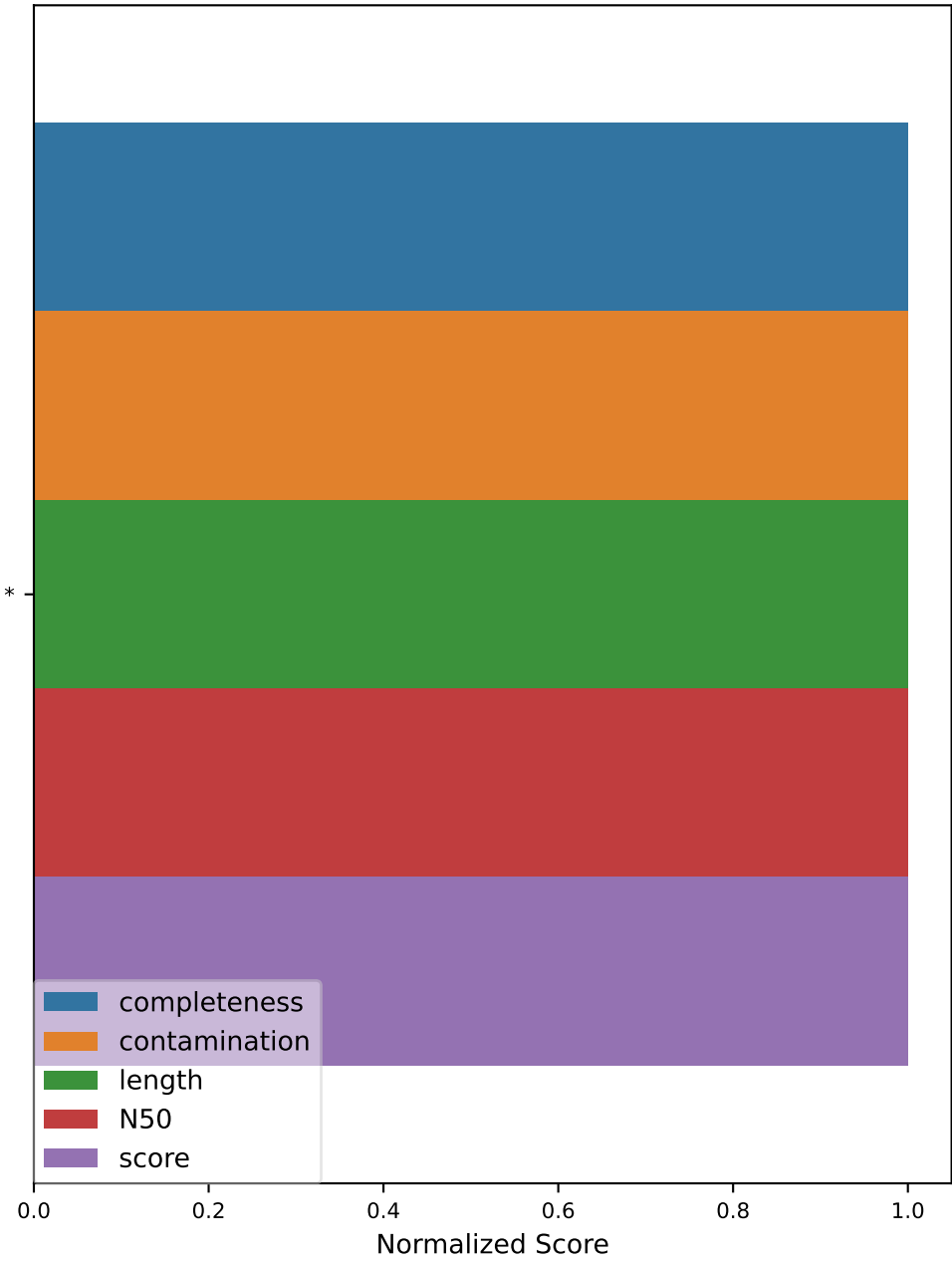
49647\_1#100\_clean\_bin.14\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 33\_0

genome

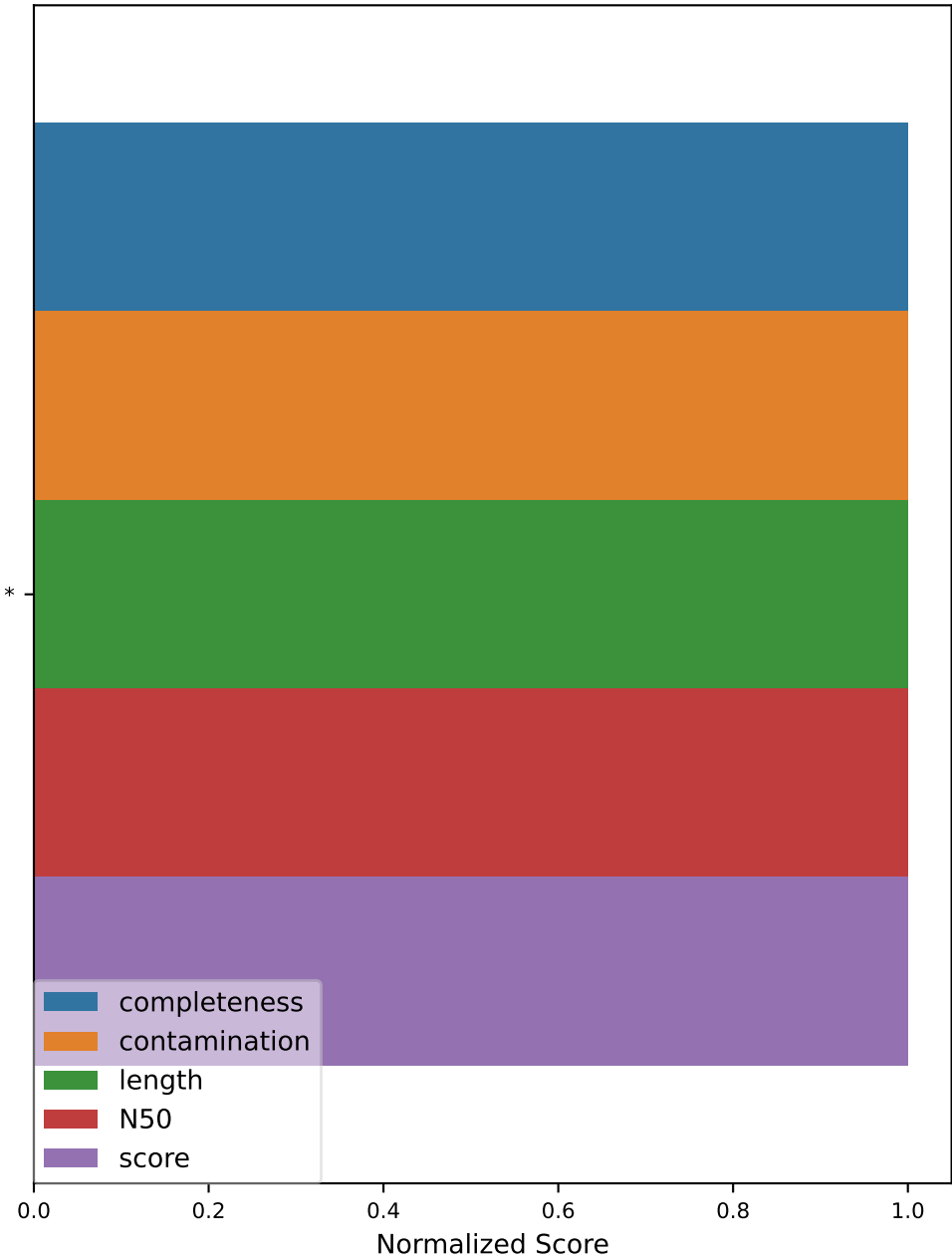
49647\_1#100\_clean\_bin.23\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 34\_0

genome

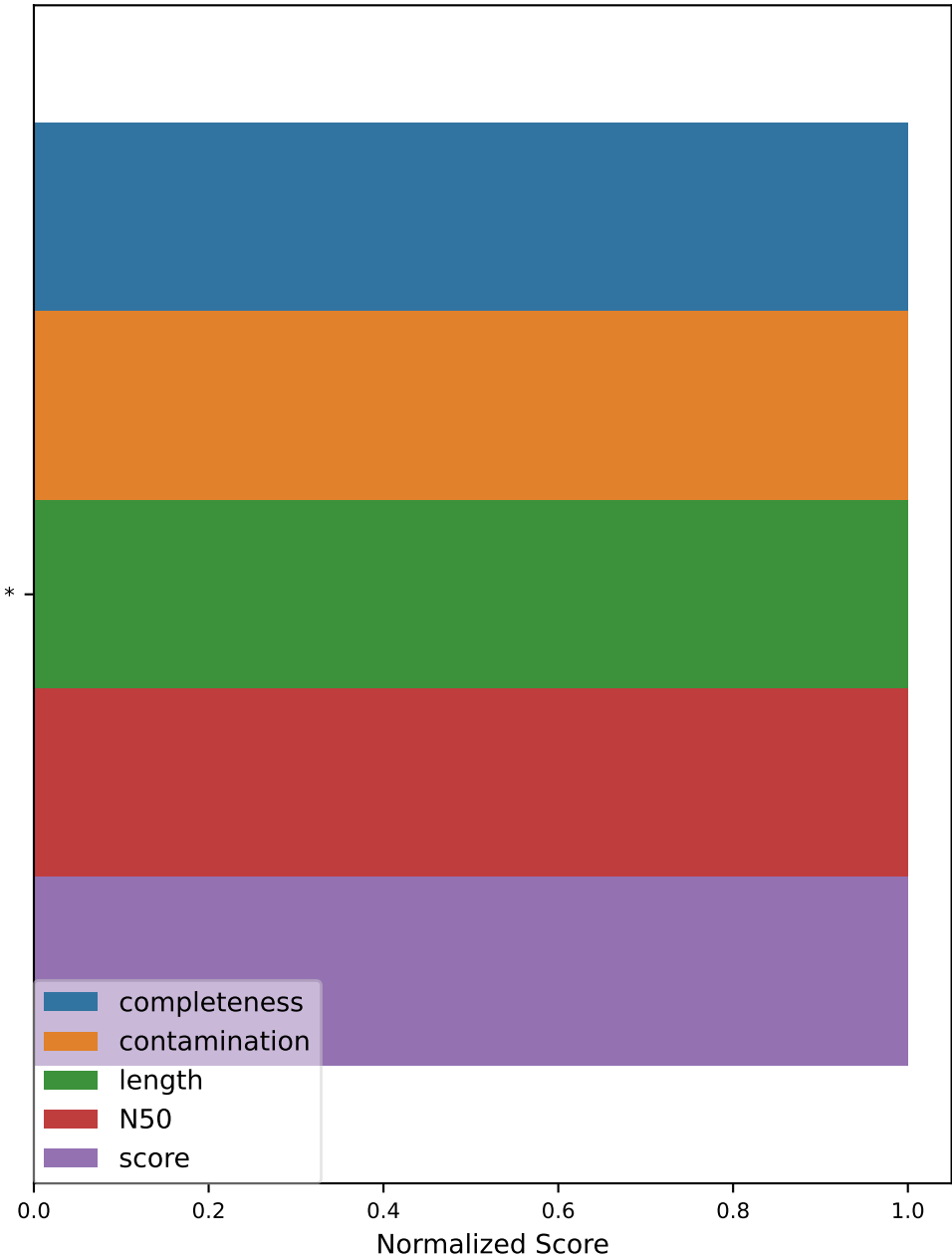
49647\_1#102\_clean\_bin.8\_4\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 35\_0

genome

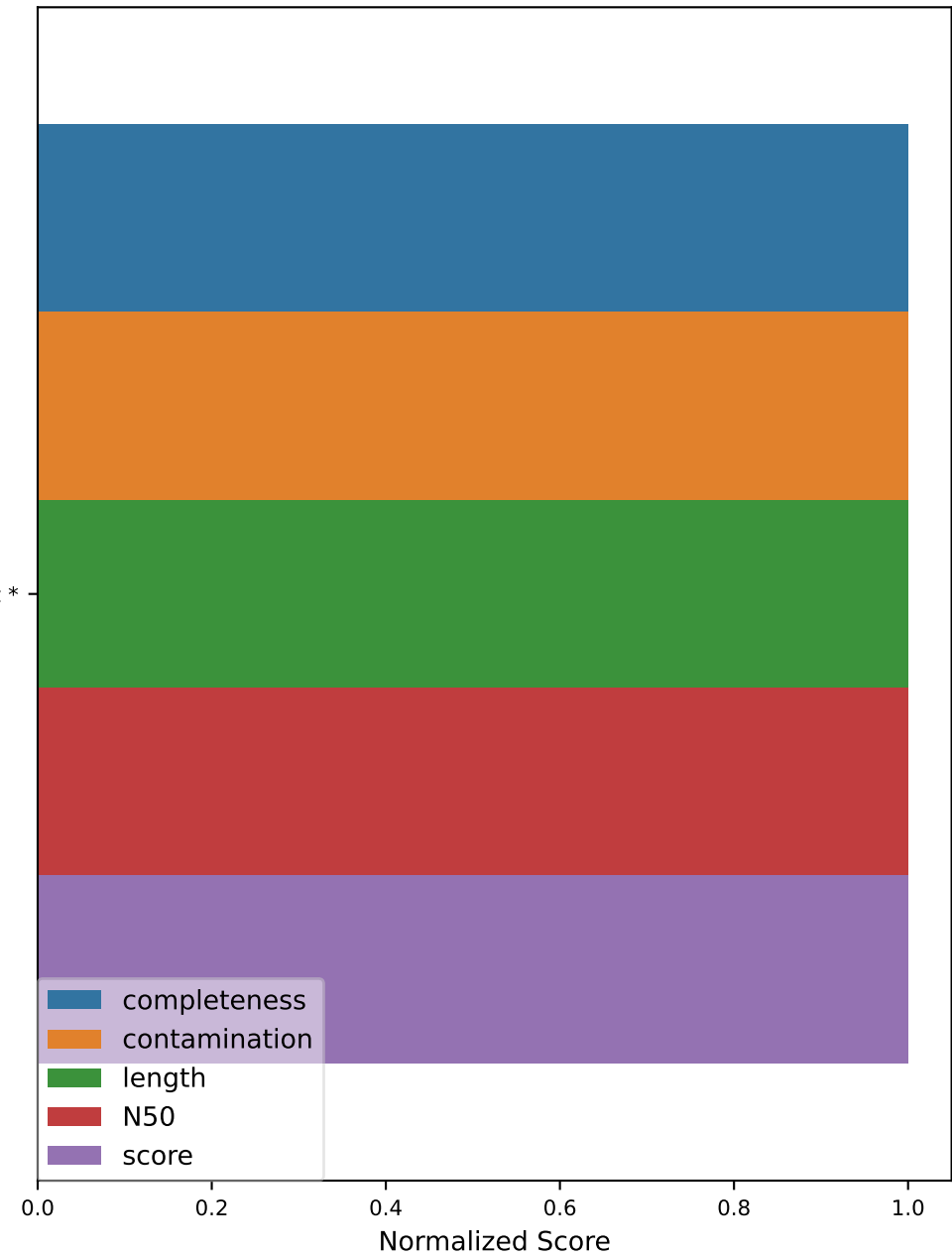
49647\_1#2\_clean\_bin.2.strict\_filtered\_kept\_contigs.fasta.gz \*



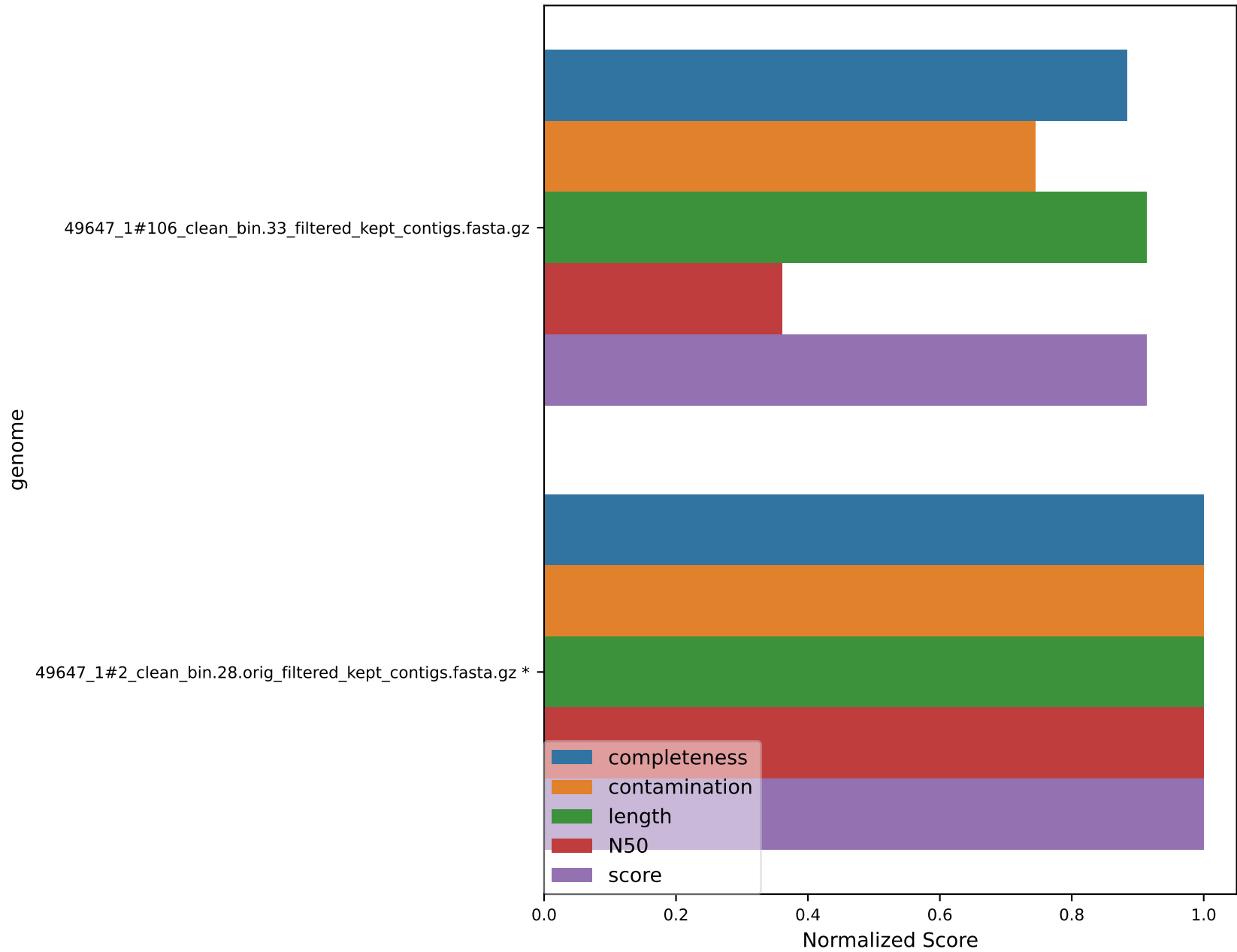
Scoring of cluster 36\_0

genome

49647\_1#70\_clean\_bin.1.strict\_filtered\_kept\_contigs.fasta.gz \*



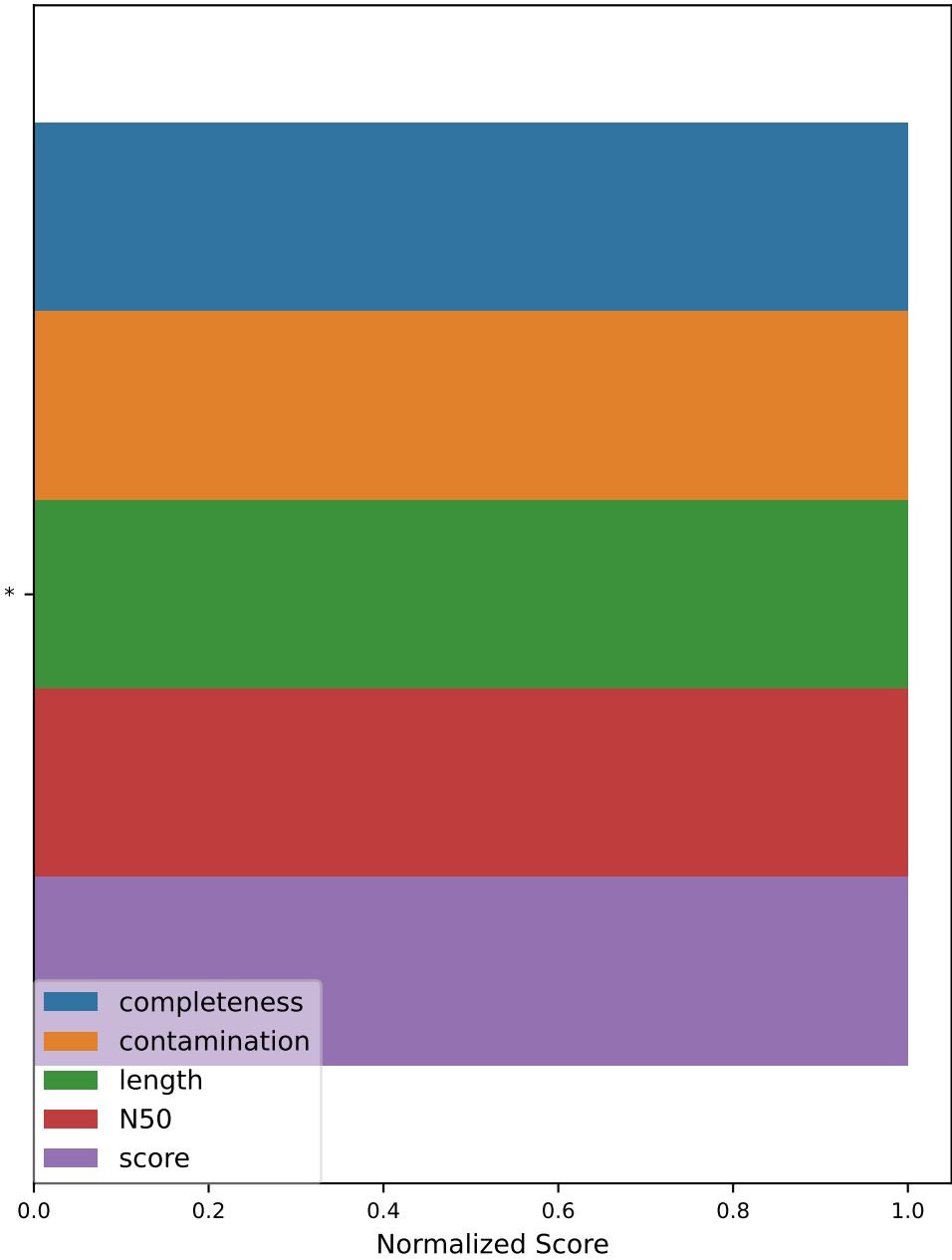
Scoring of cluster 37\_1



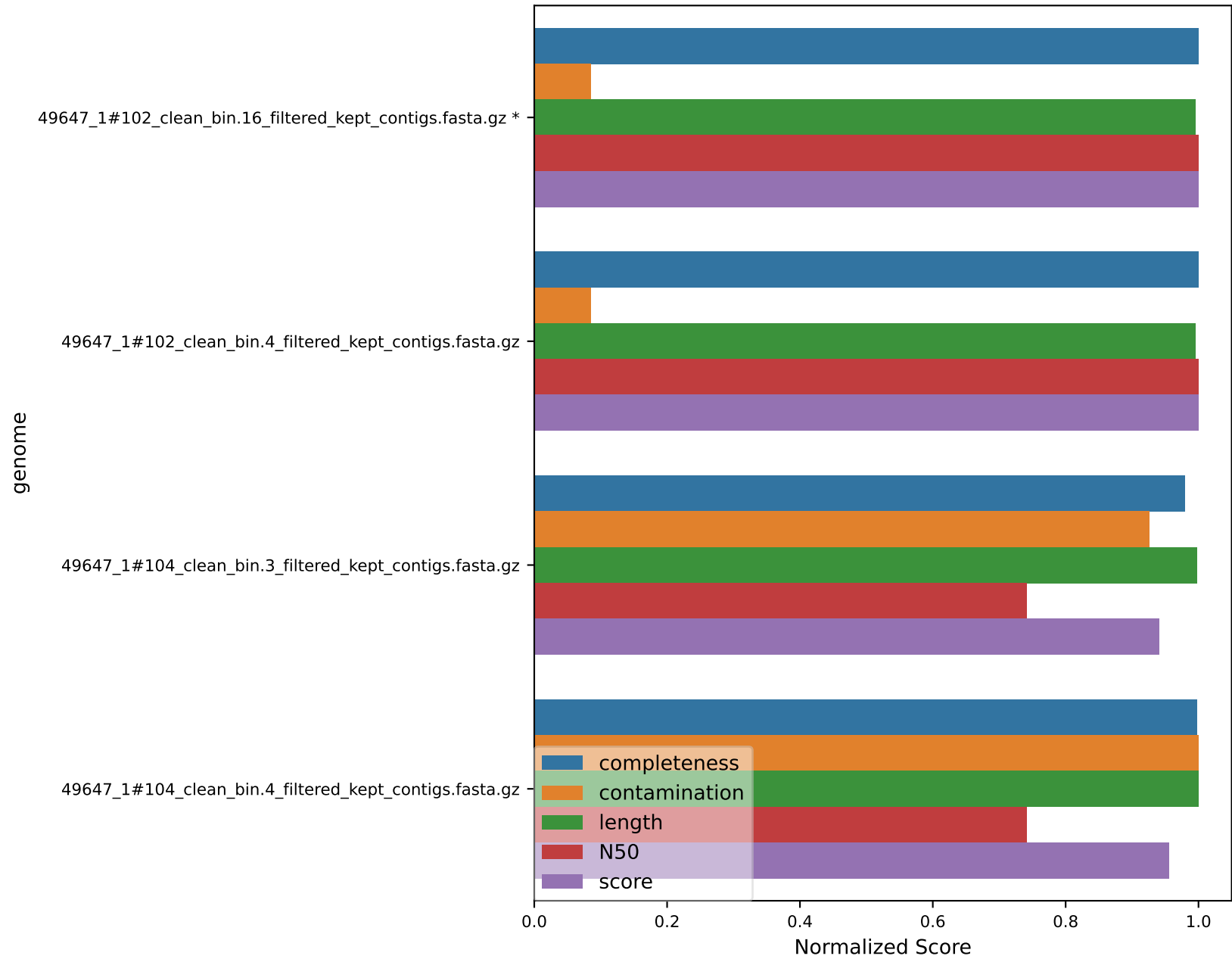
Scoring of cluster 38\_0

genome

49647\_1#106\_clean\_bin.31\_21\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 39\_1

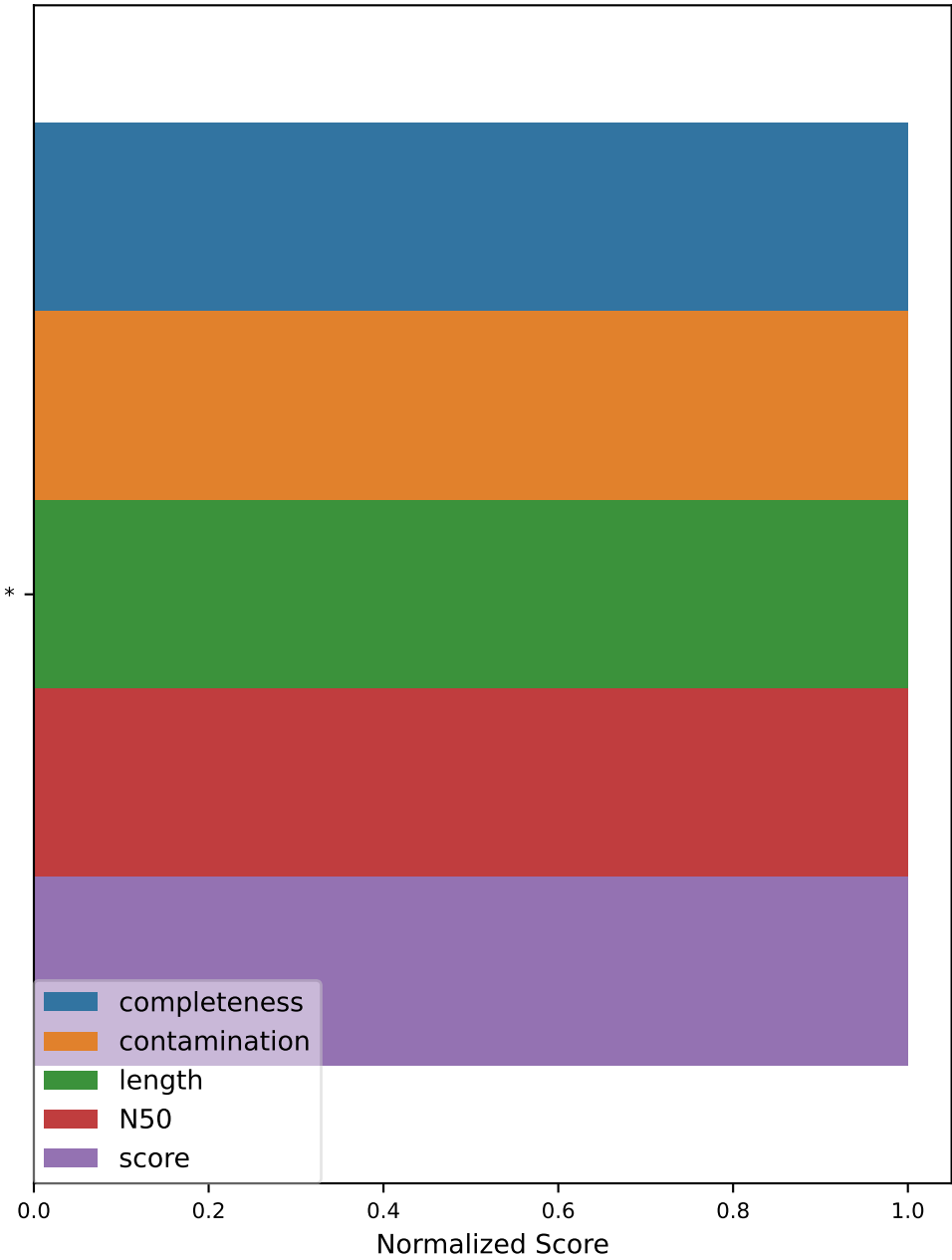




Scoring of cluster 39\_2

genome

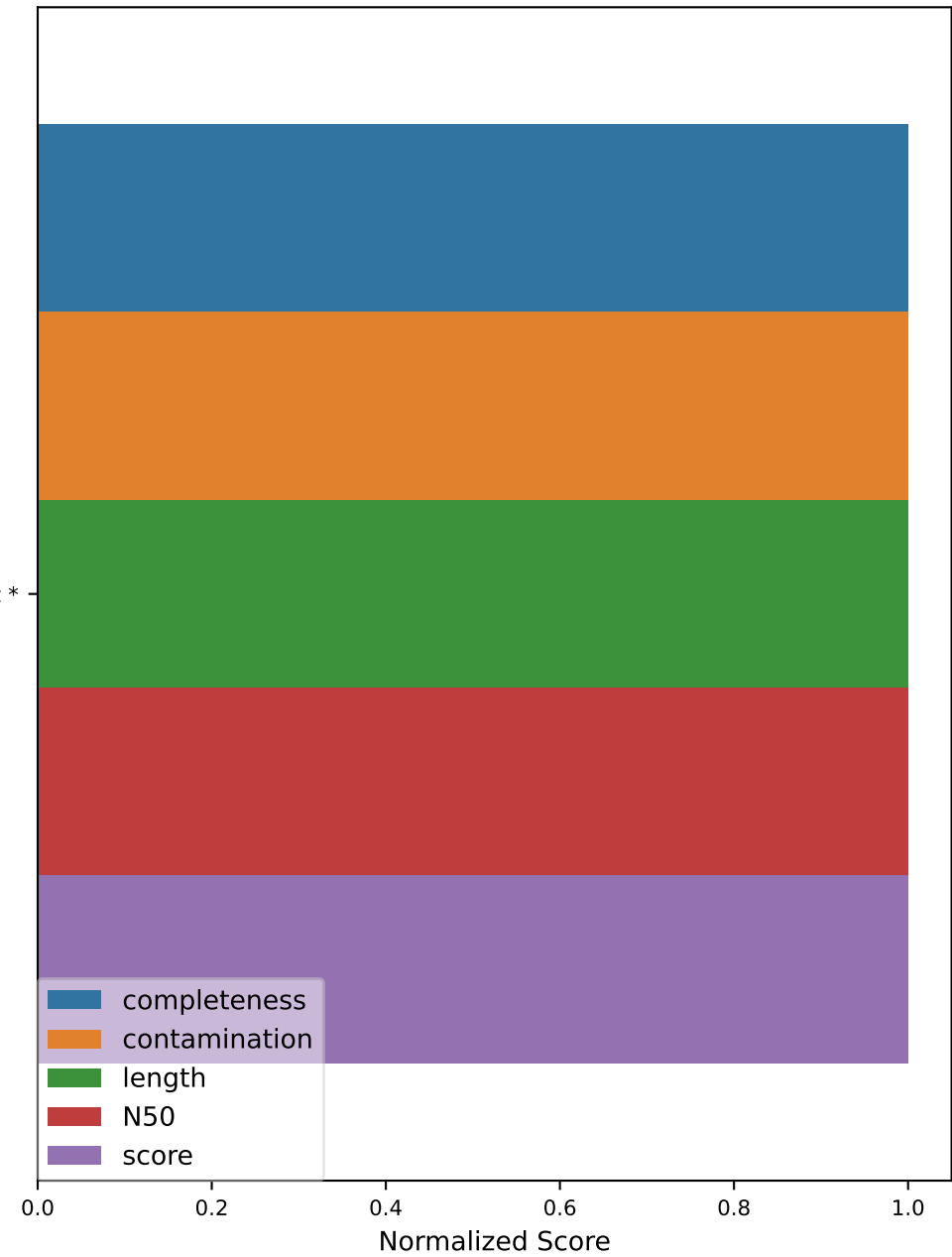
49647\_1#100\_clean\_bin.26\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 40\_0

genome

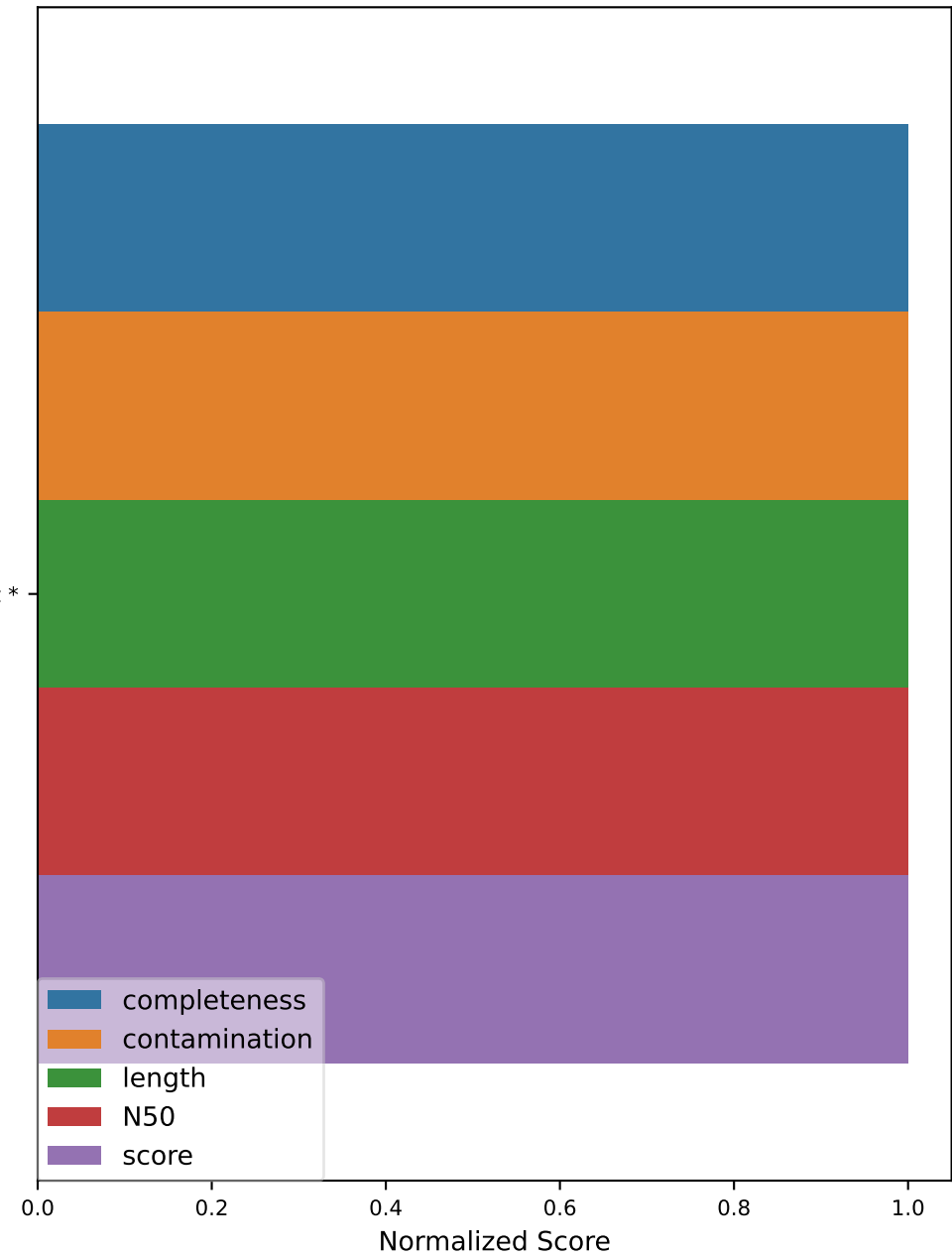
49647\_1#100\_clean\_bin.3\_filtered\_kept\_contigs.fasta.gz \*



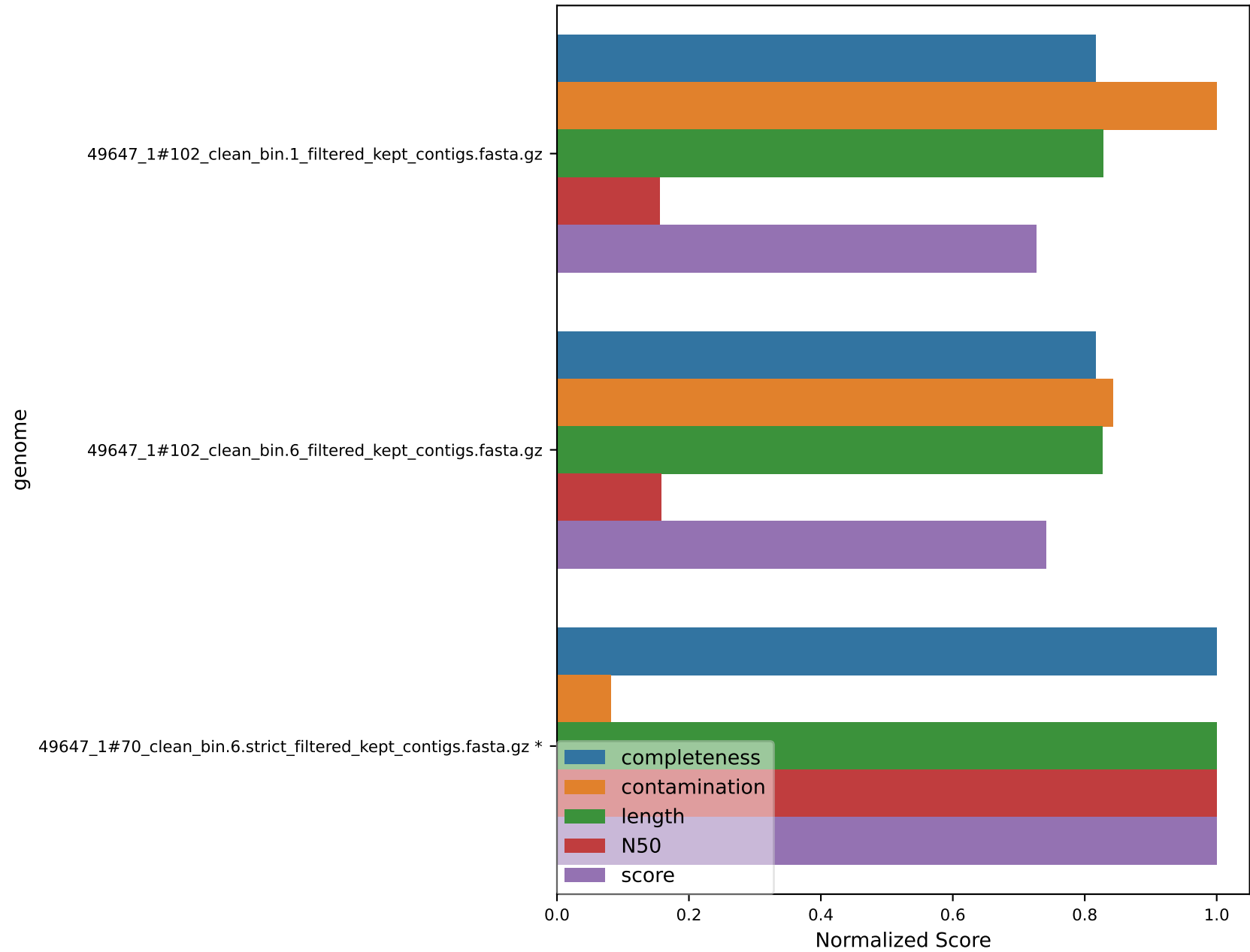
Scoring of cluster 41\_0

genome

49647\_1#106\_clean\_bin.17\_1\_filtered\_kept\_contigs.fasta.gz \*



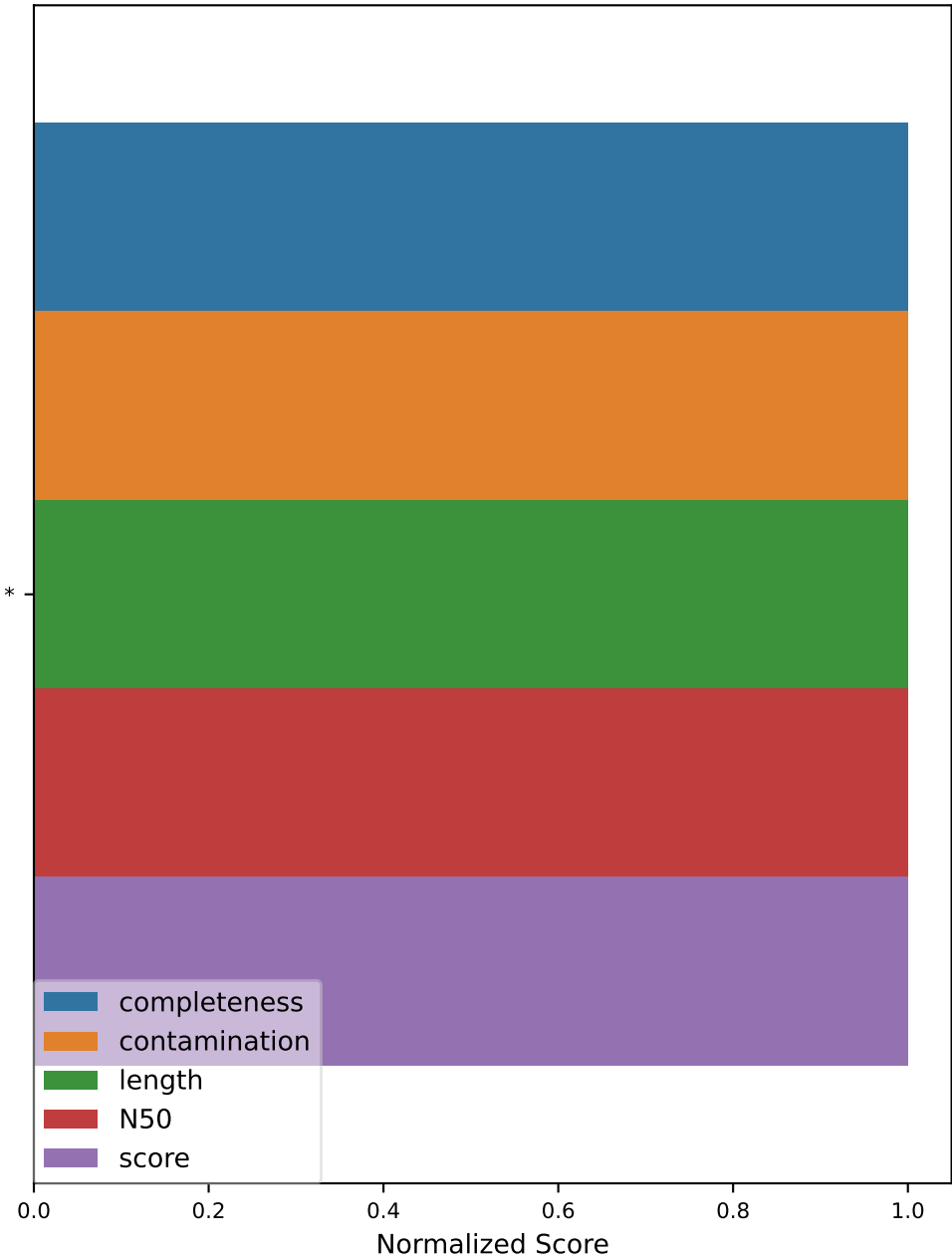
Scoring of cluster 42\_1



Scoring of cluster 43\_0

genome

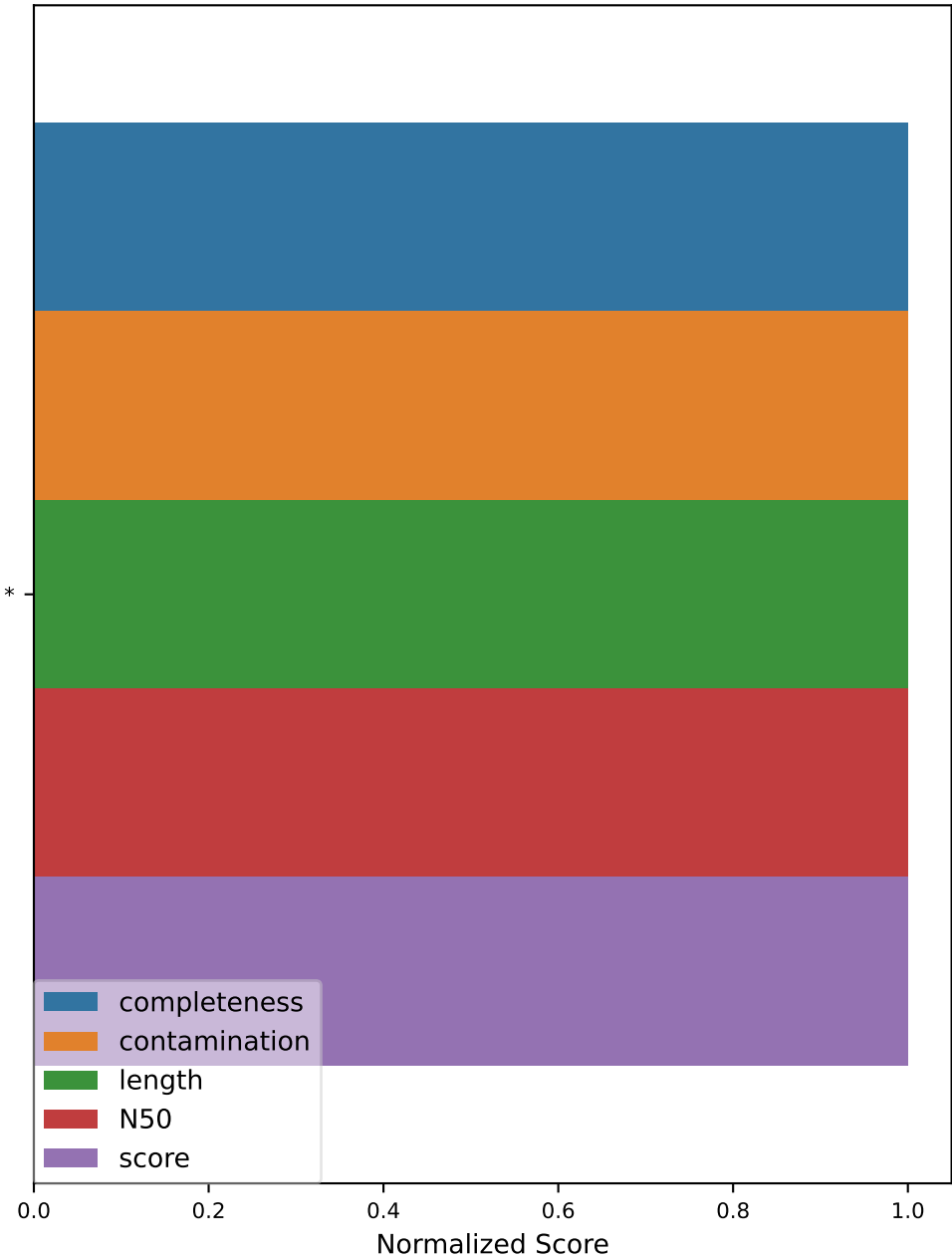
49647\_1#100\_clean\_bin.11\_filtered\_kept\_contigs.fasta.gz \*



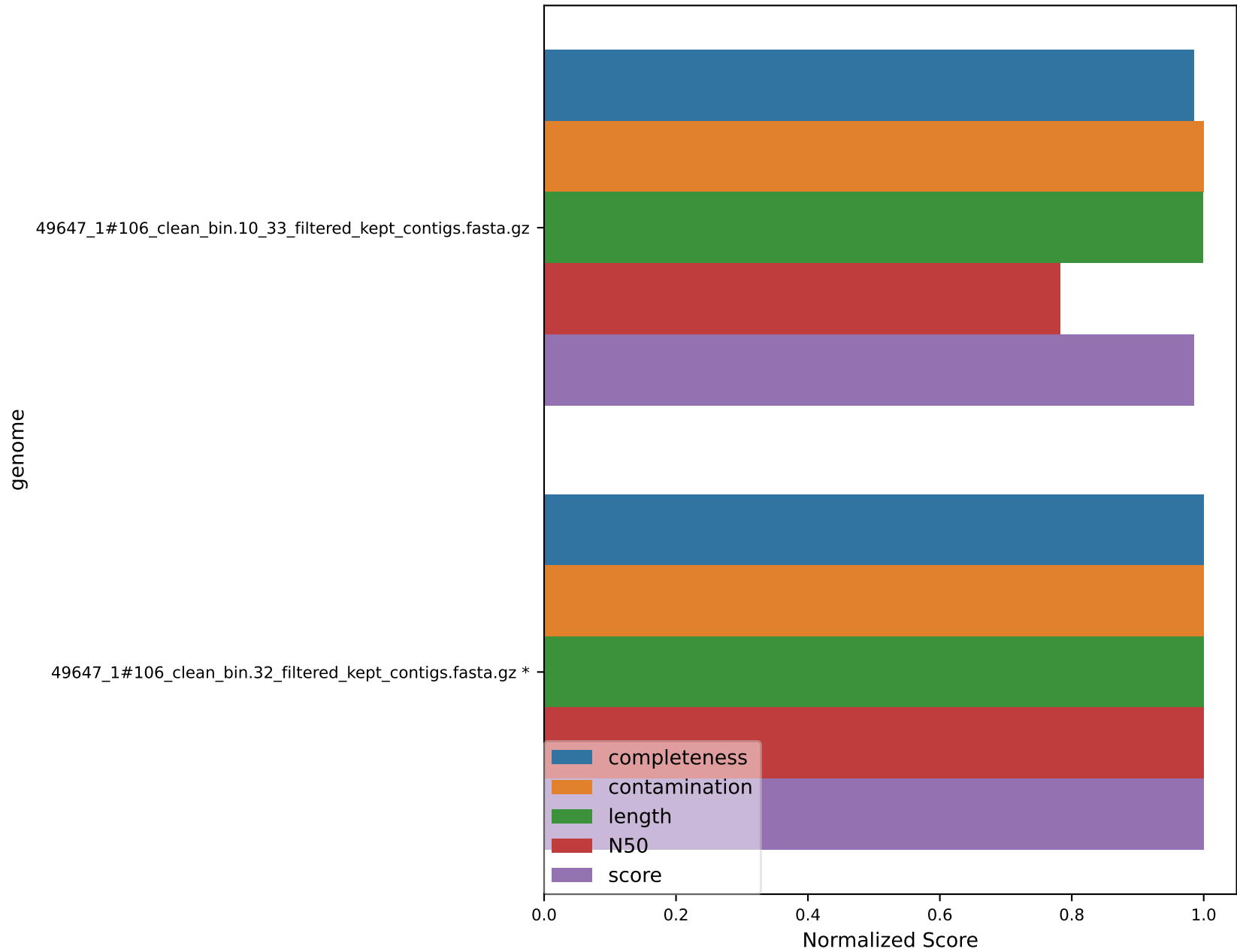
Scoring of cluster 44\_0

genome

49647\_1#106\_clean\_bin.12\_23\_filtered\_kept\_contigs.fasta.gz \*



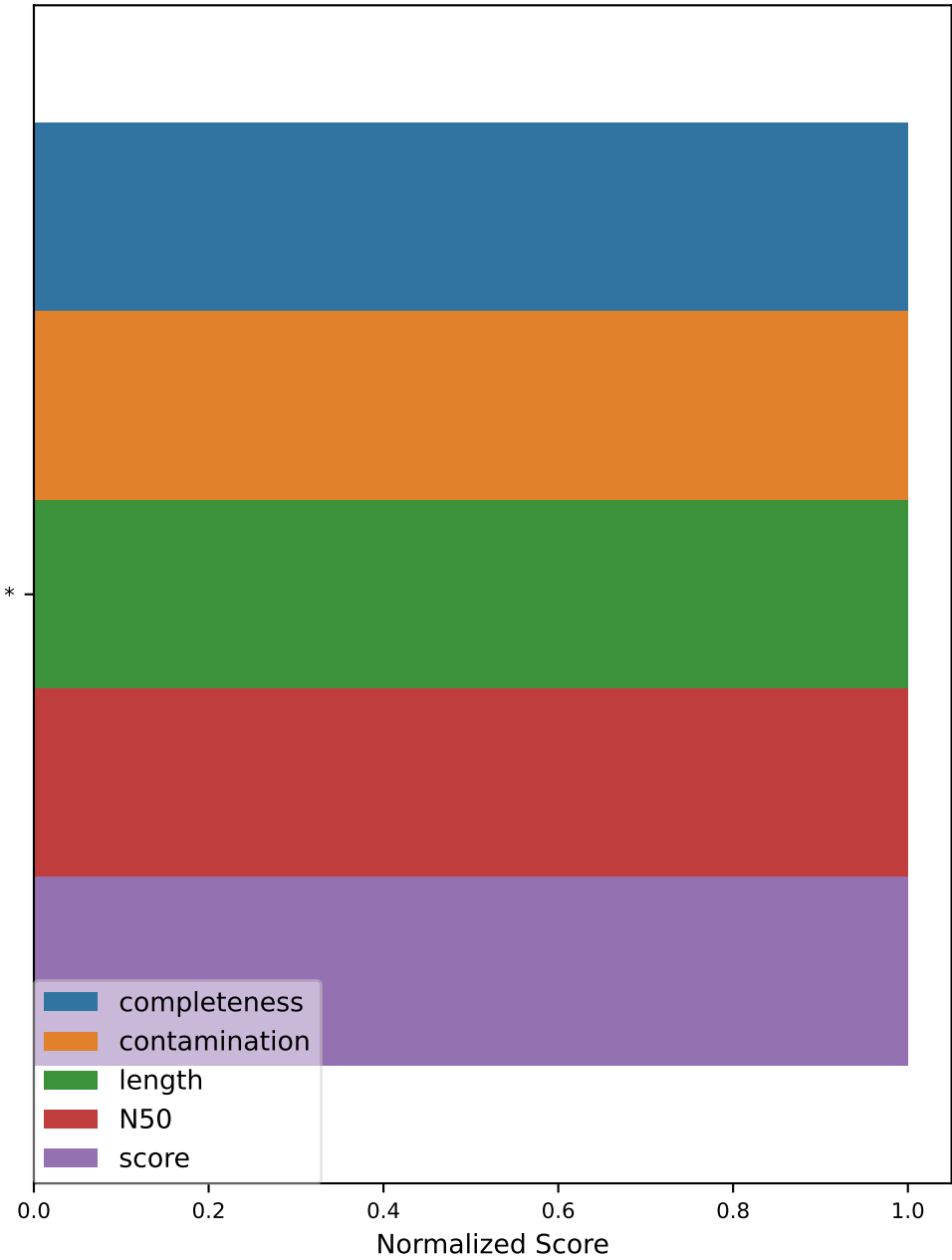
Scoring of cluster 45\_1



Scoring of cluster 46\_0

genome

49647\_1#2\_clean\_bin.10.permissive\_filtered\_kept\_contigs.fasta.gz \*

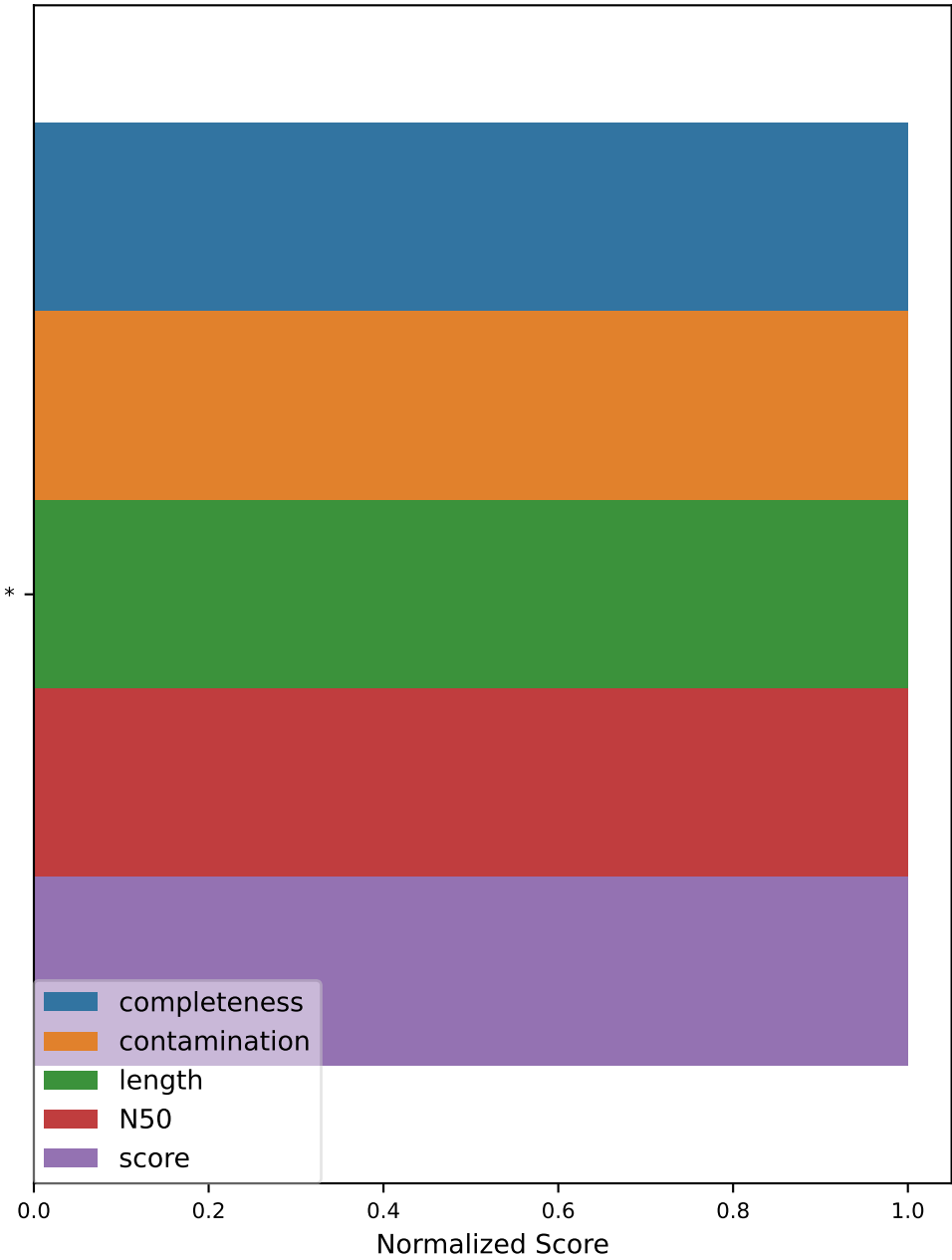




Scoring of cluster 47\_0

genome

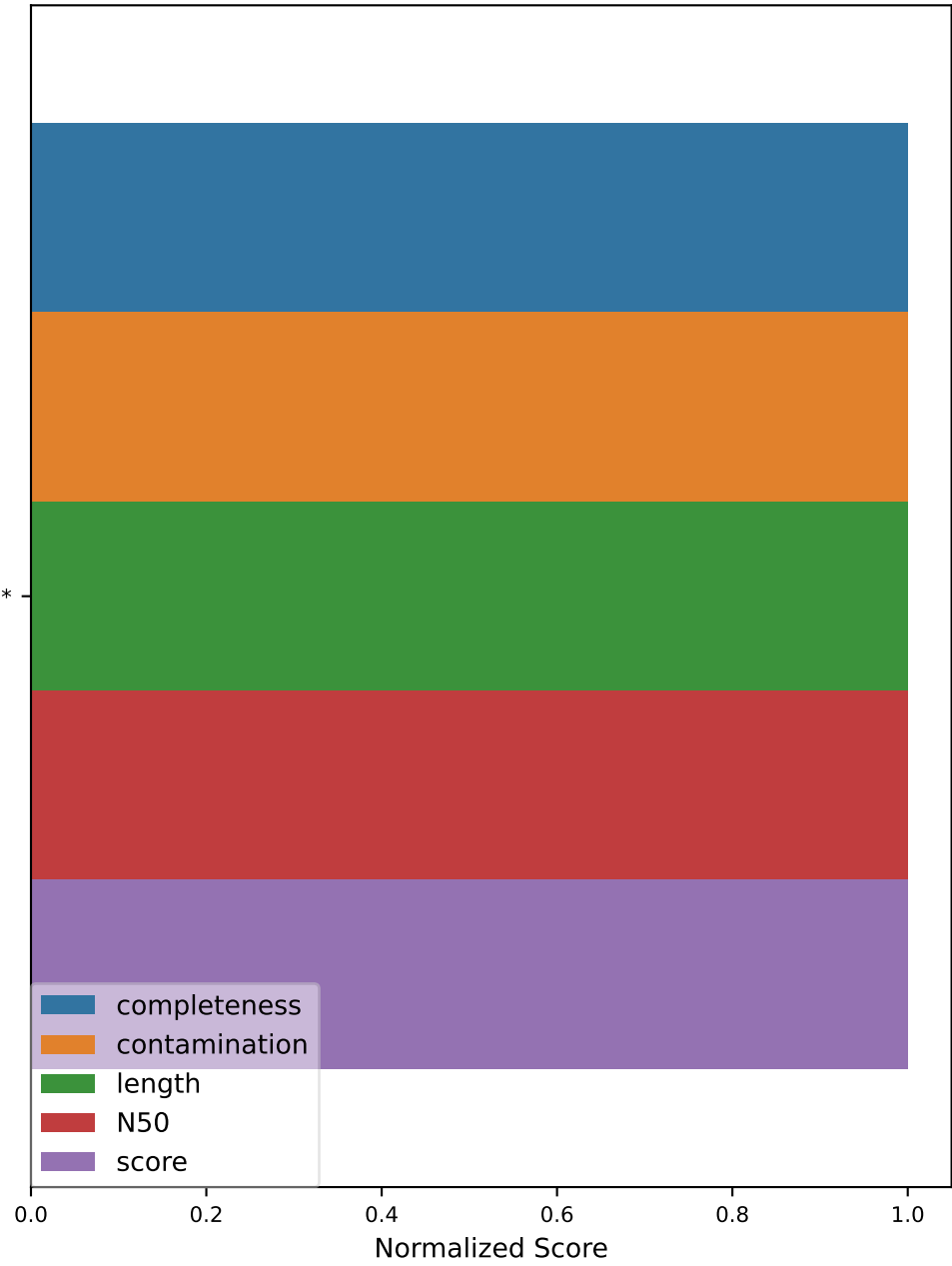
49647\_1#106\_clean\_bin.37\_filtered\_kept\_contigs.fasta.gz \*



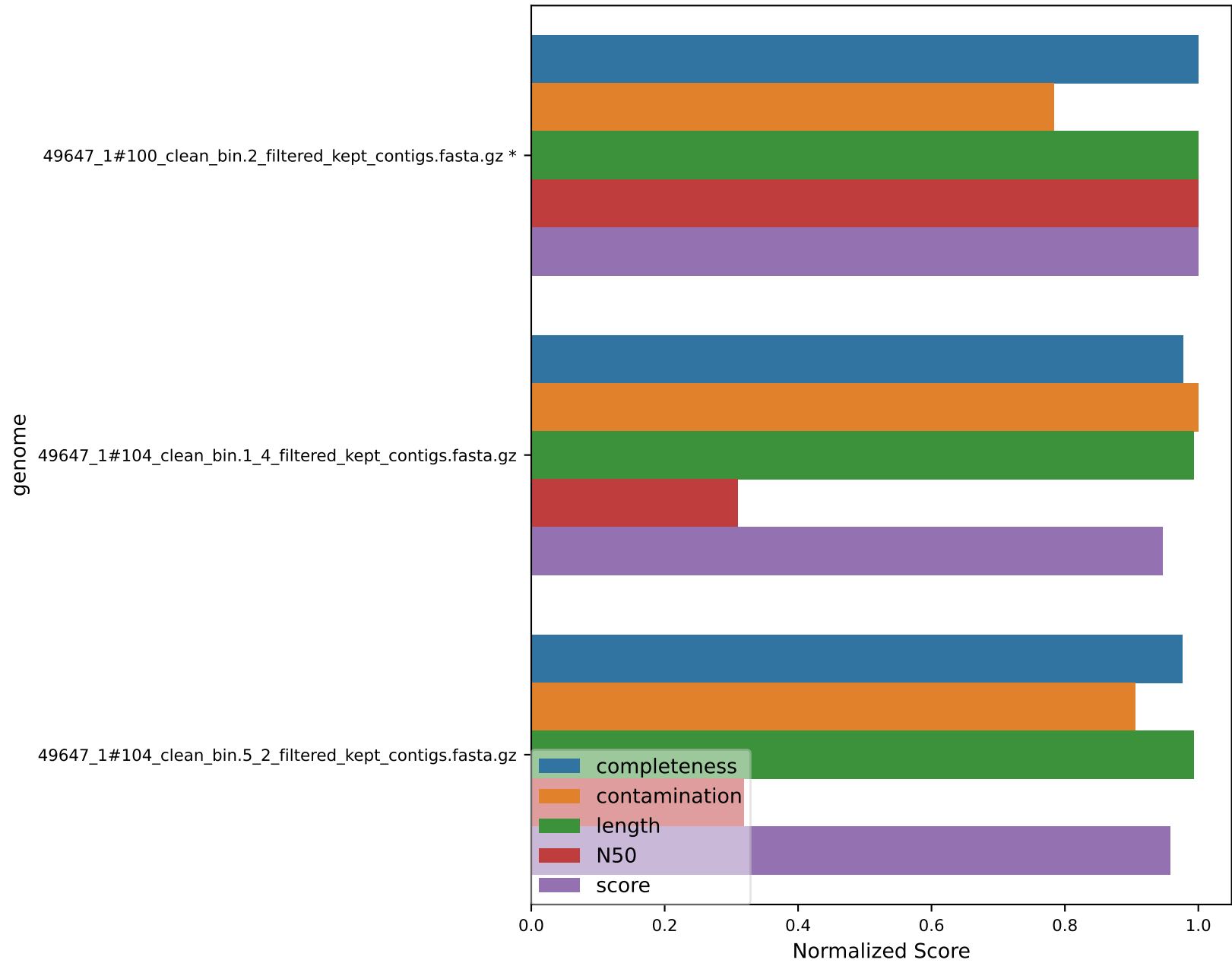
Scoring of cluster 48\_0

genome

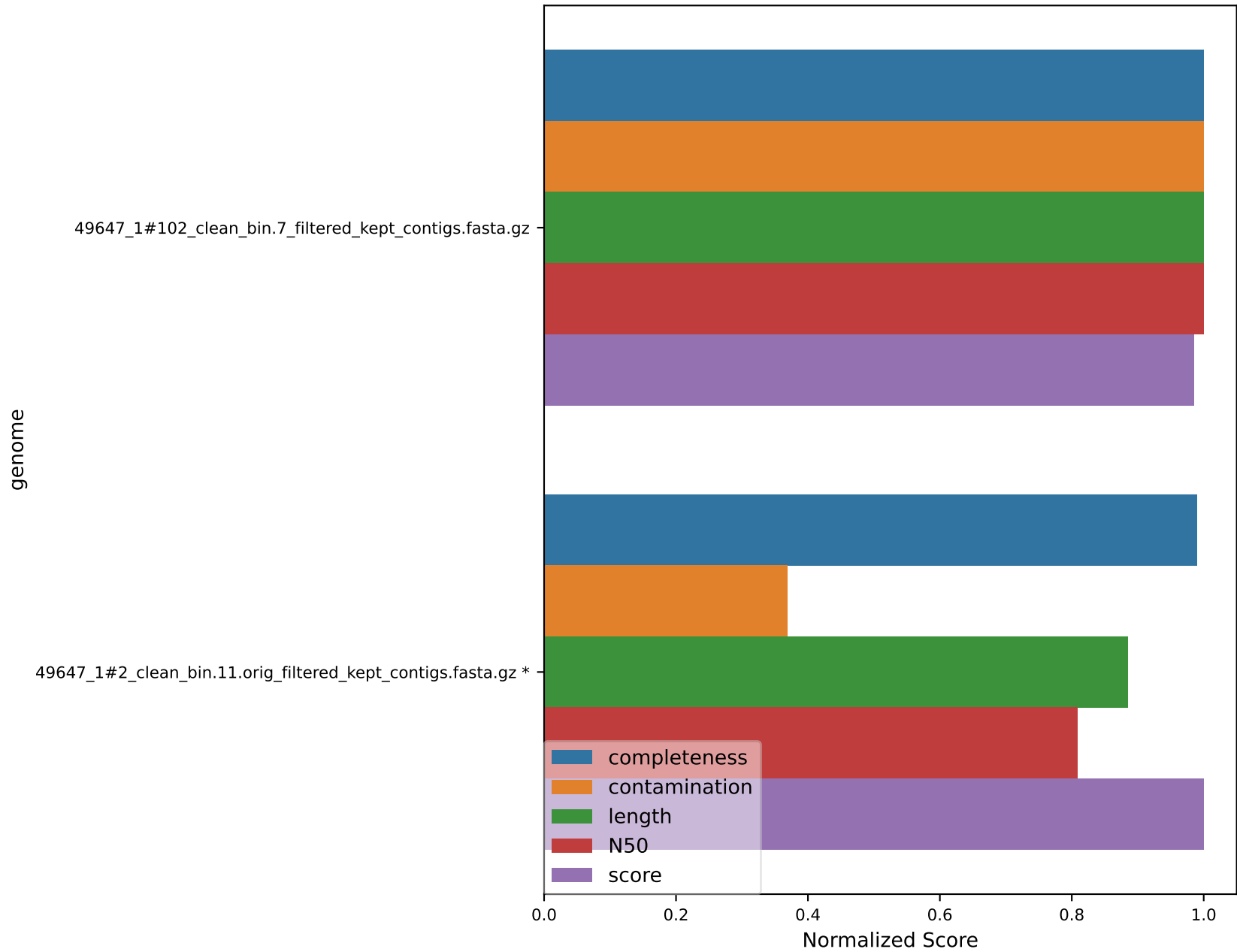
49647\_1#106\_clean\_bin.26\_filtered\_kept\_contigs.fasta.gz \*



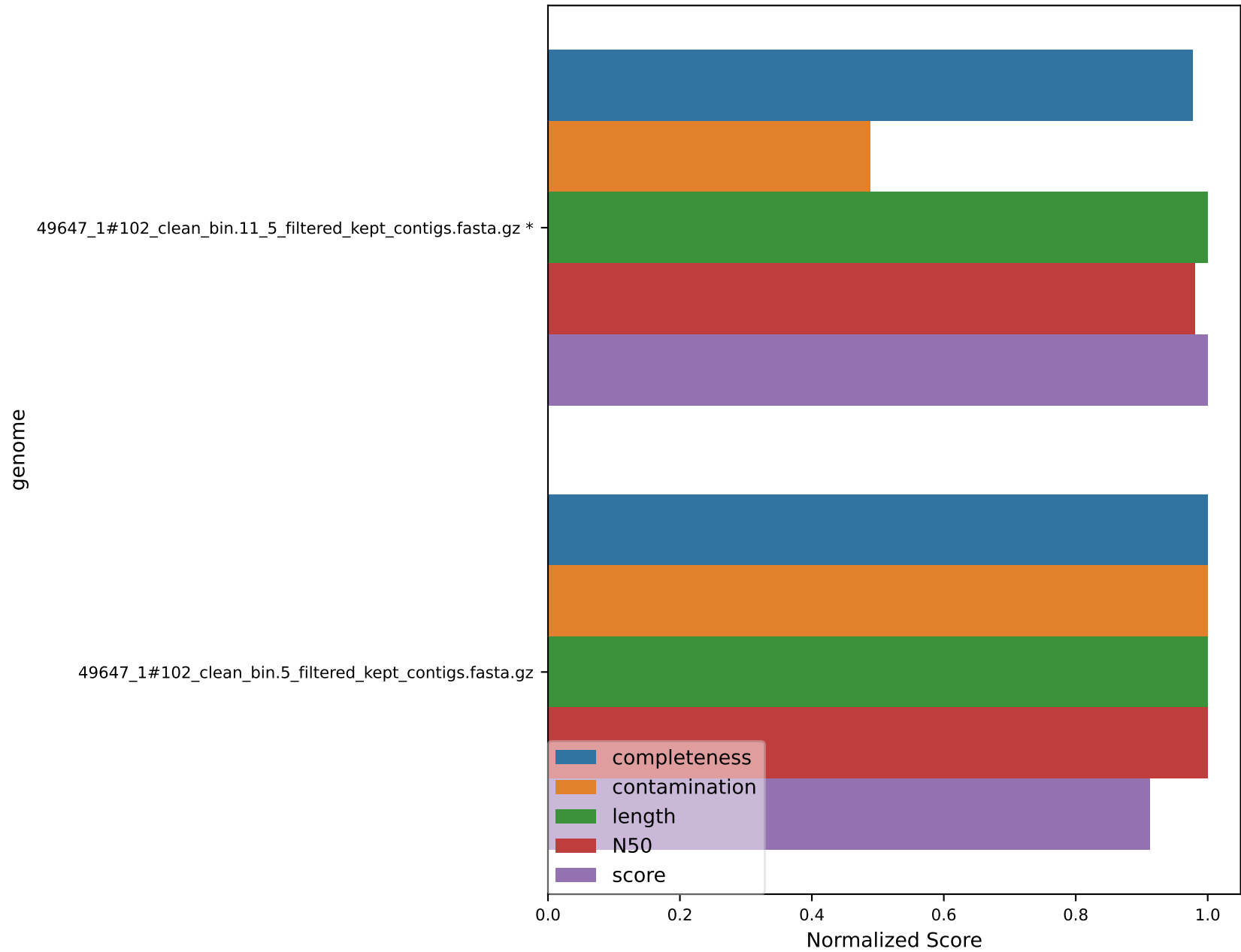
Scoring of cluster 49\_1



Scoring of cluster 50\_1



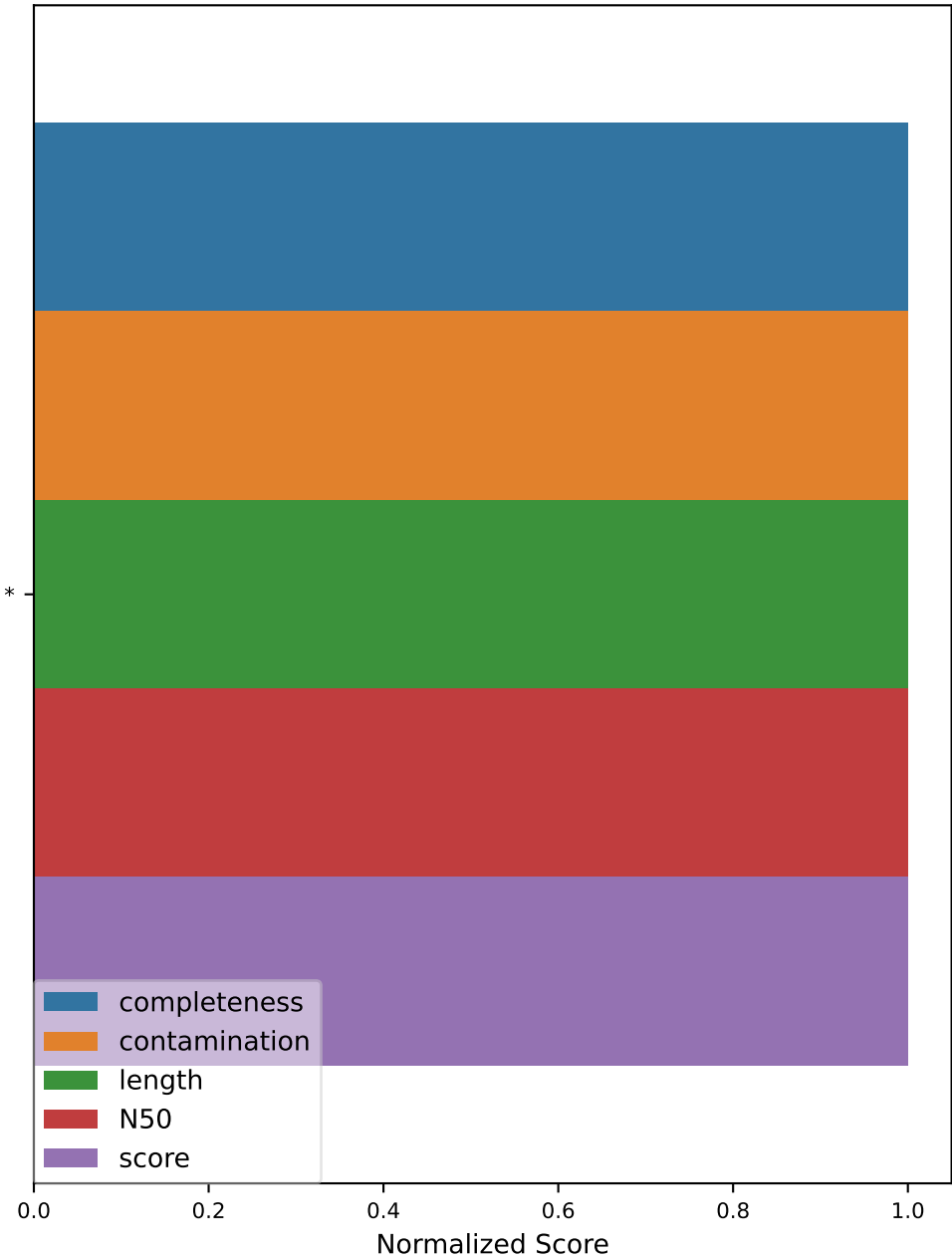
Scoring of cluster 51\_1



Scoring of cluster 52\_0

genome

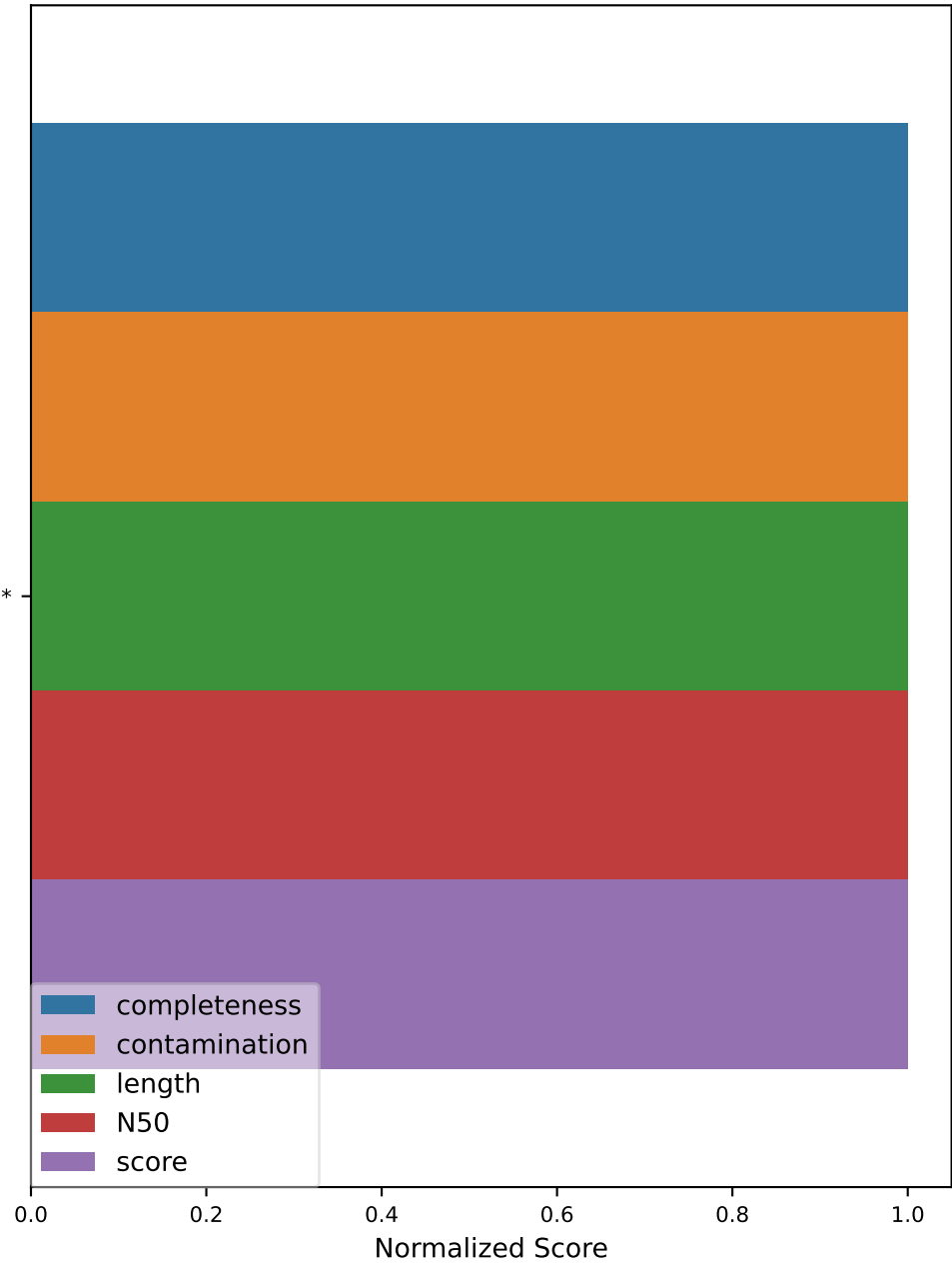
49647\_1#2\_clean\_bin.16.orig\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 53\_0

genome

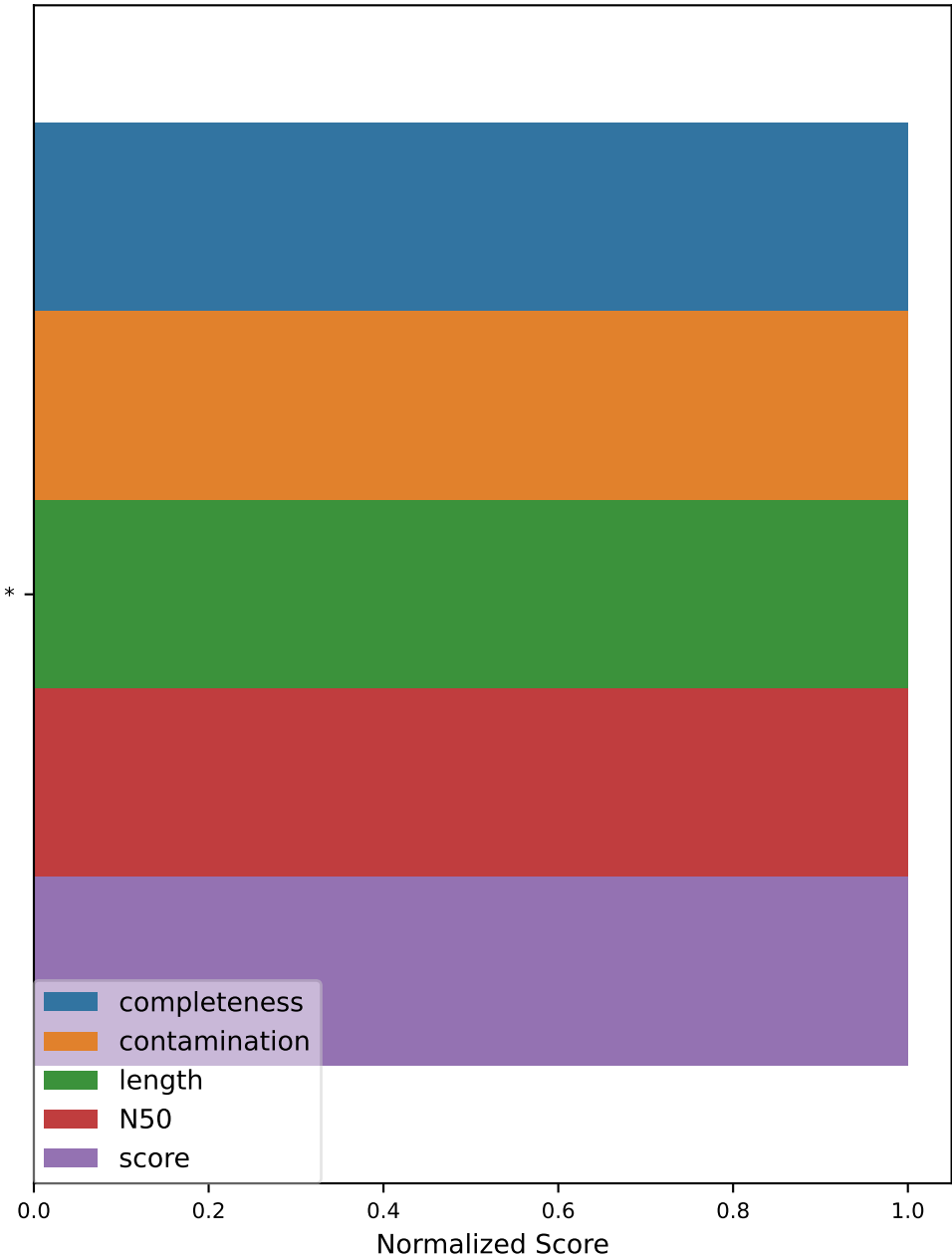
49647\_1#106\_clean\_bin.35\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 54\_0

genome

49647\_1#106\_clean\_bin.25\_filtered\_kept\_contigs.fasta.gz \*

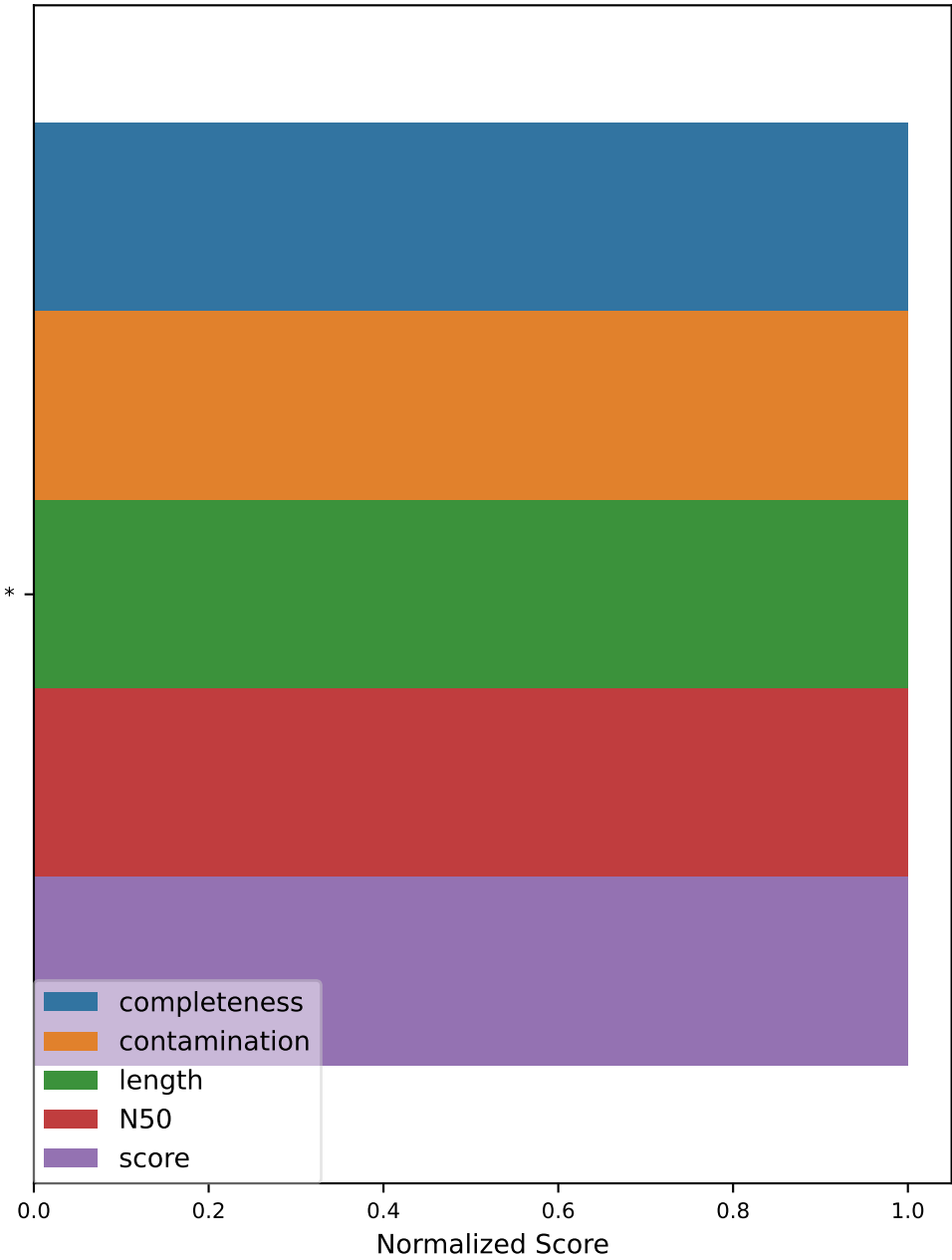




Scoring of cluster 55\_0

genome

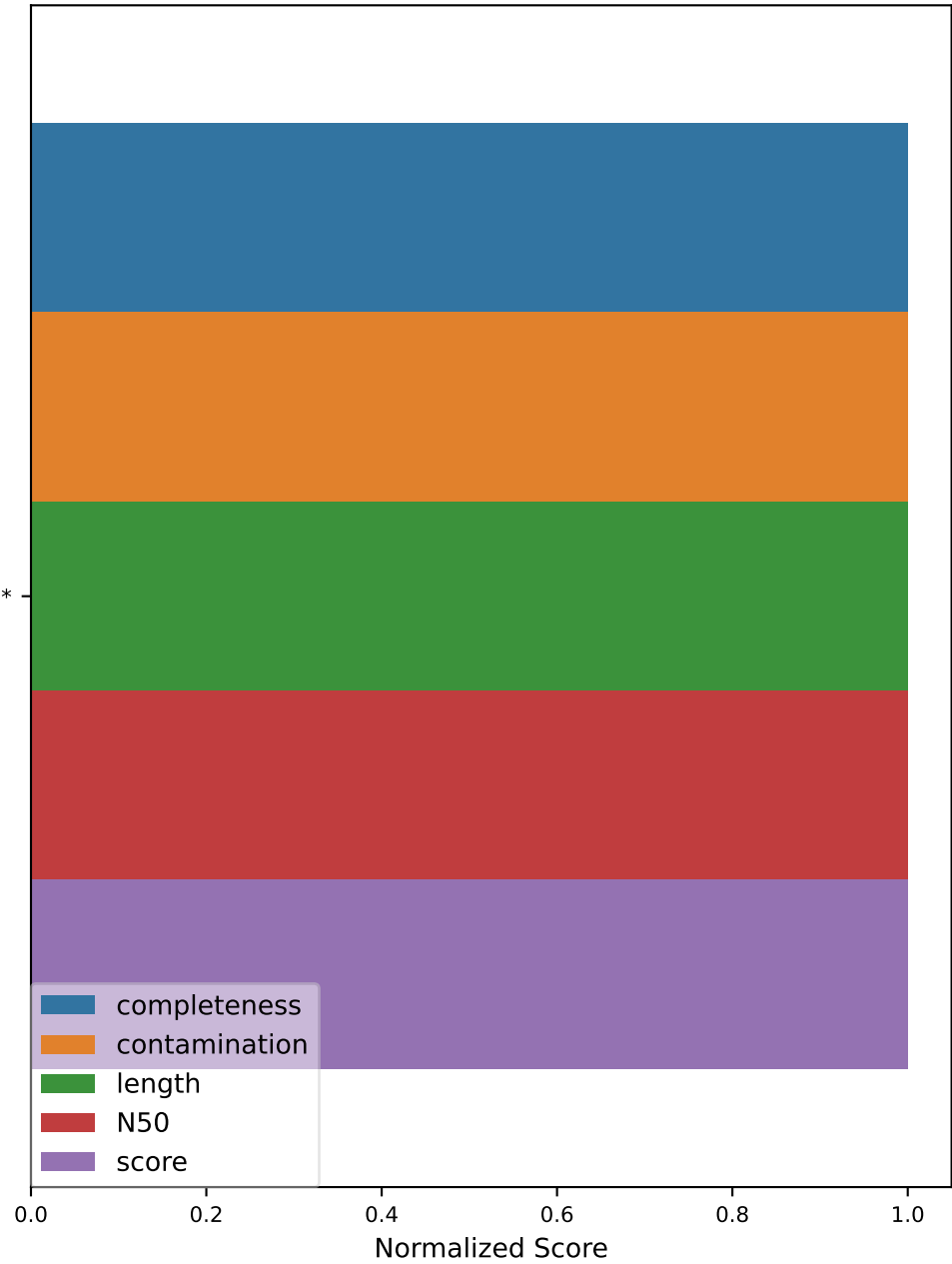
49647\_1#106\_clean\_bin.28\_31\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 56\_0

genome

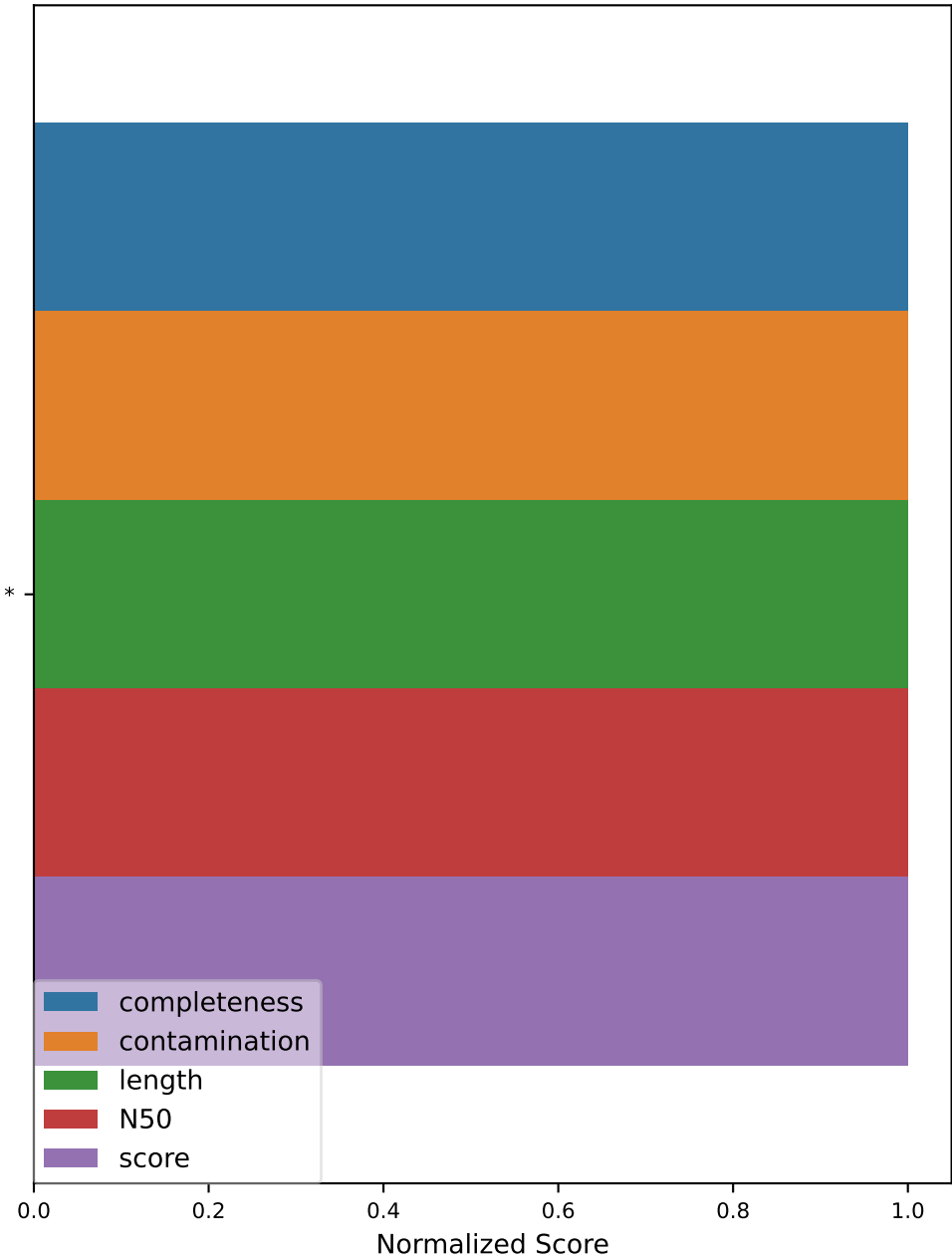
49647\_1#106\_clean\_bin.18\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 57\_0

genome

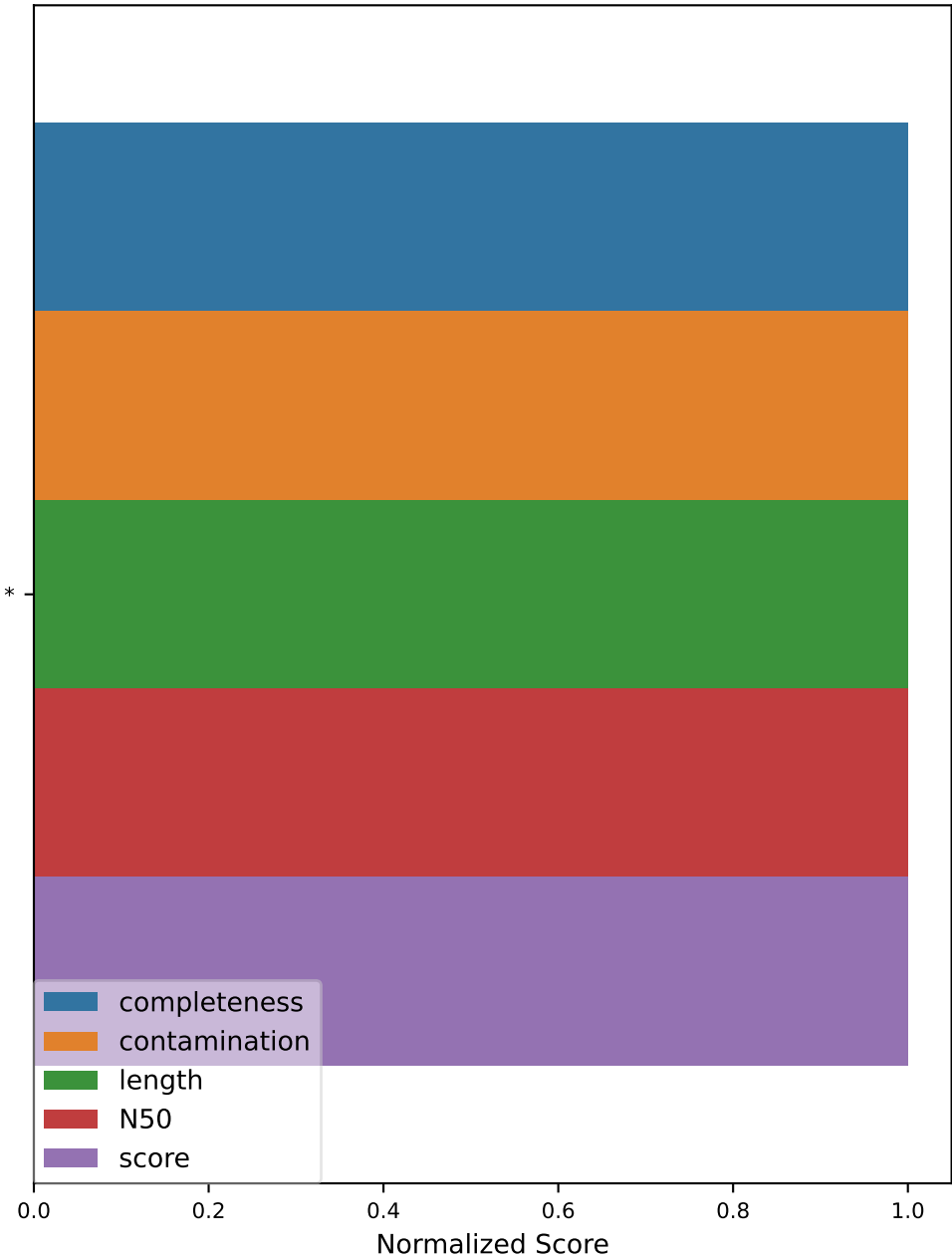
49647\_1#106\_clean\_bin.34\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 58\_1

genome

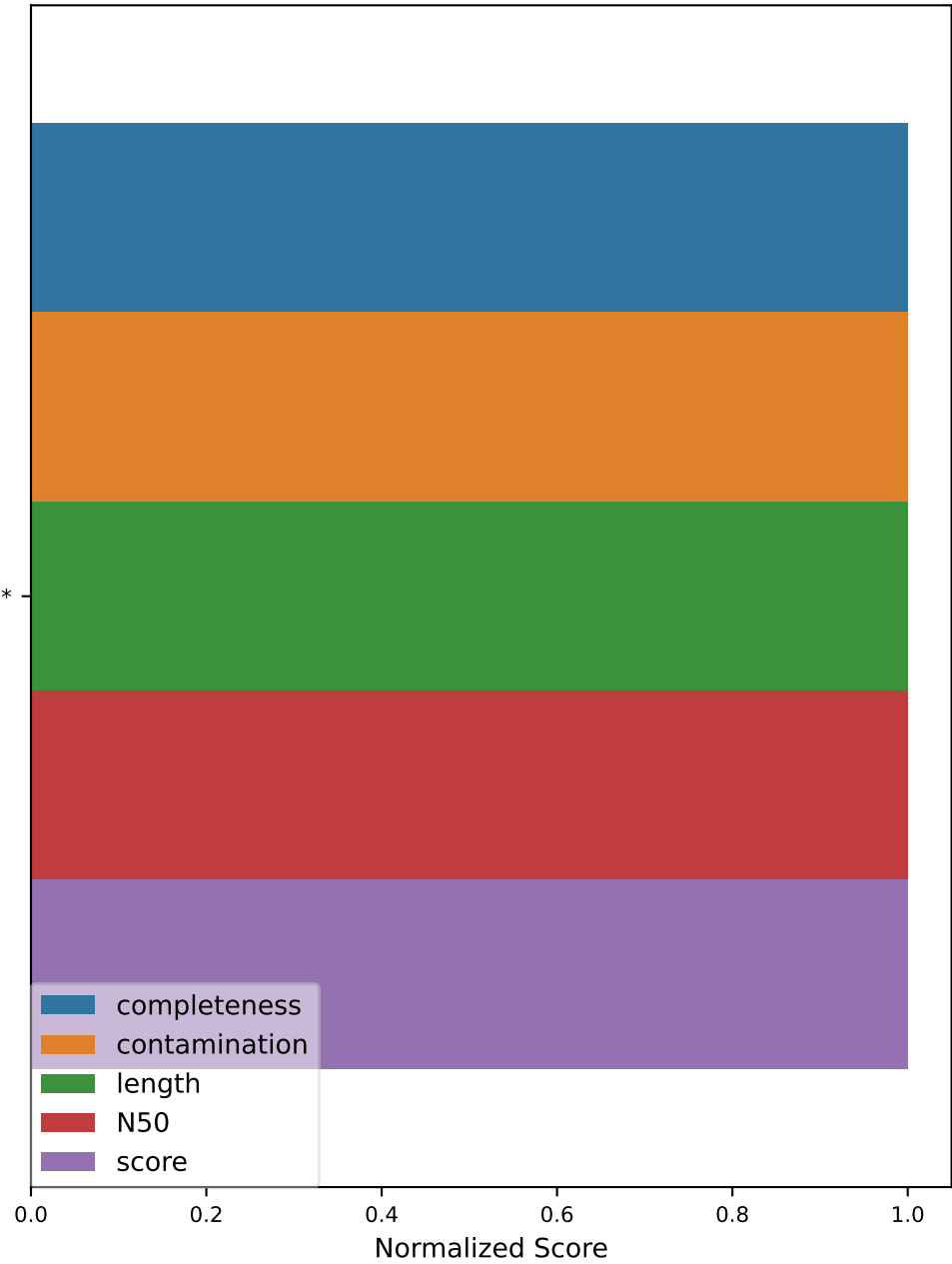
49647\_1#106\_clean\_bin.36\_6\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 58\_2

genome

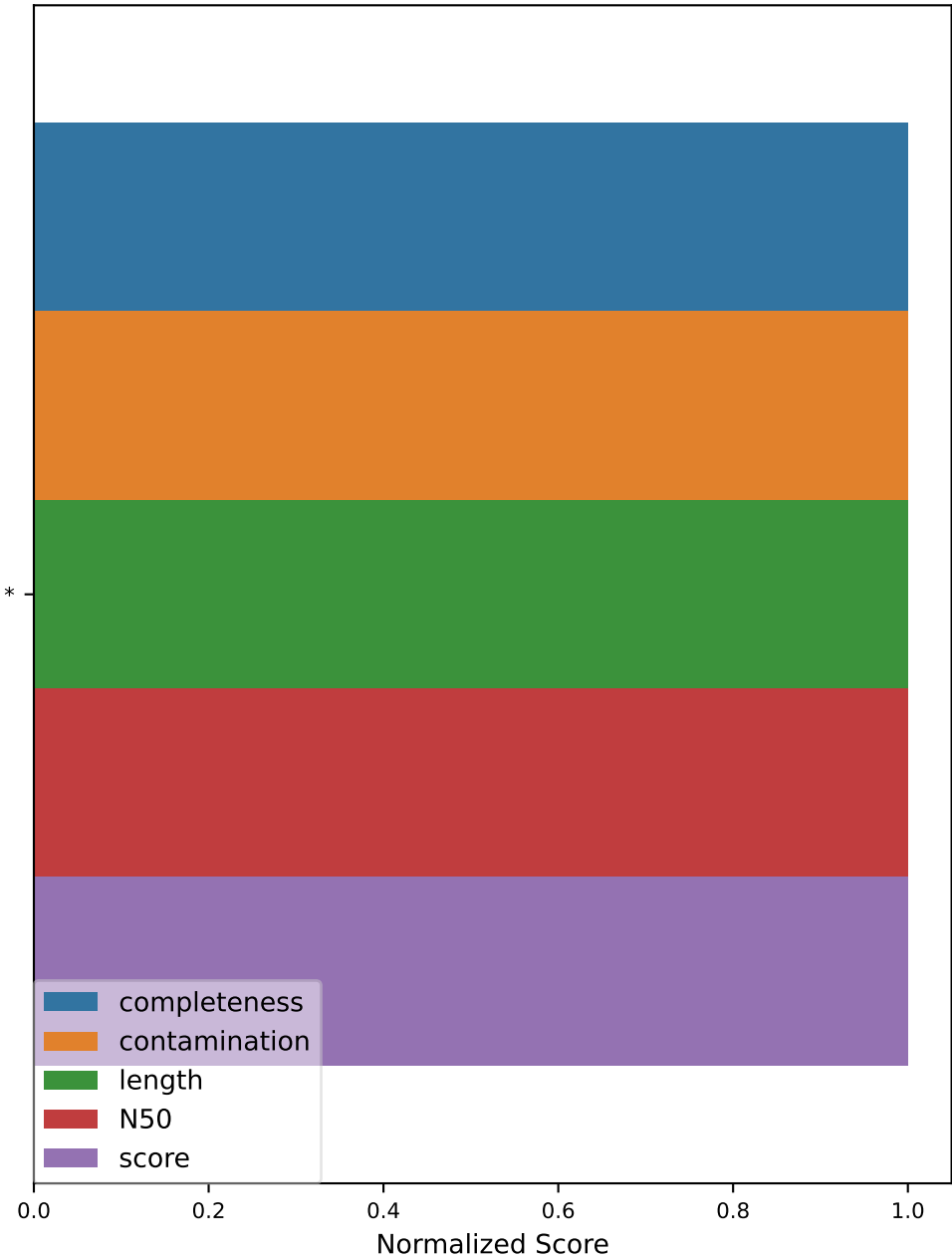
49647\_1#2\_clean\_bin.22.orig\_filtered\_kept\_contigs.fasta.gz \*



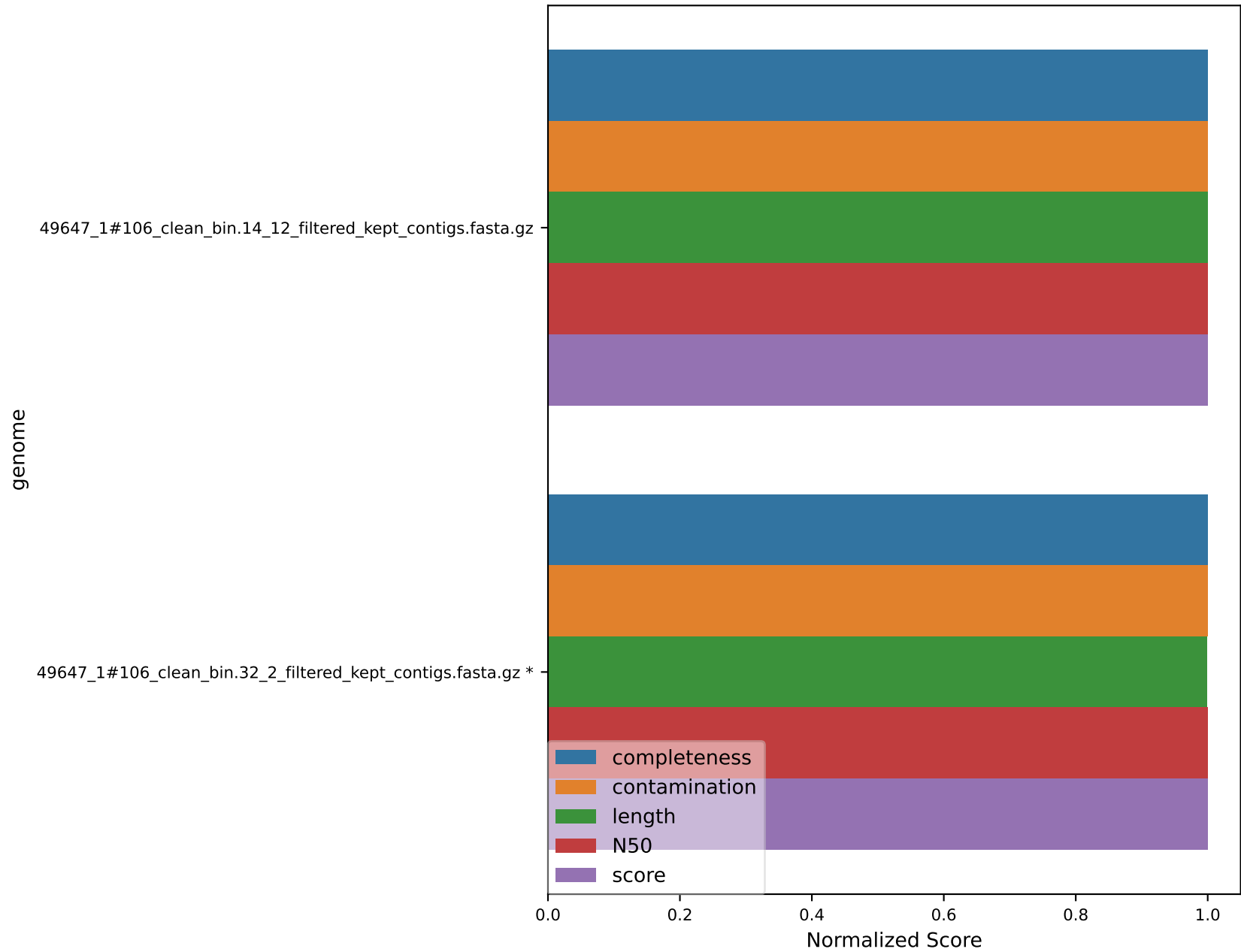
Scoring of cluster 59\_0

genome

49647\_1#106\_clean\_bin.13\_29\_filtered\_kept\_contigs.fasta.gz \*



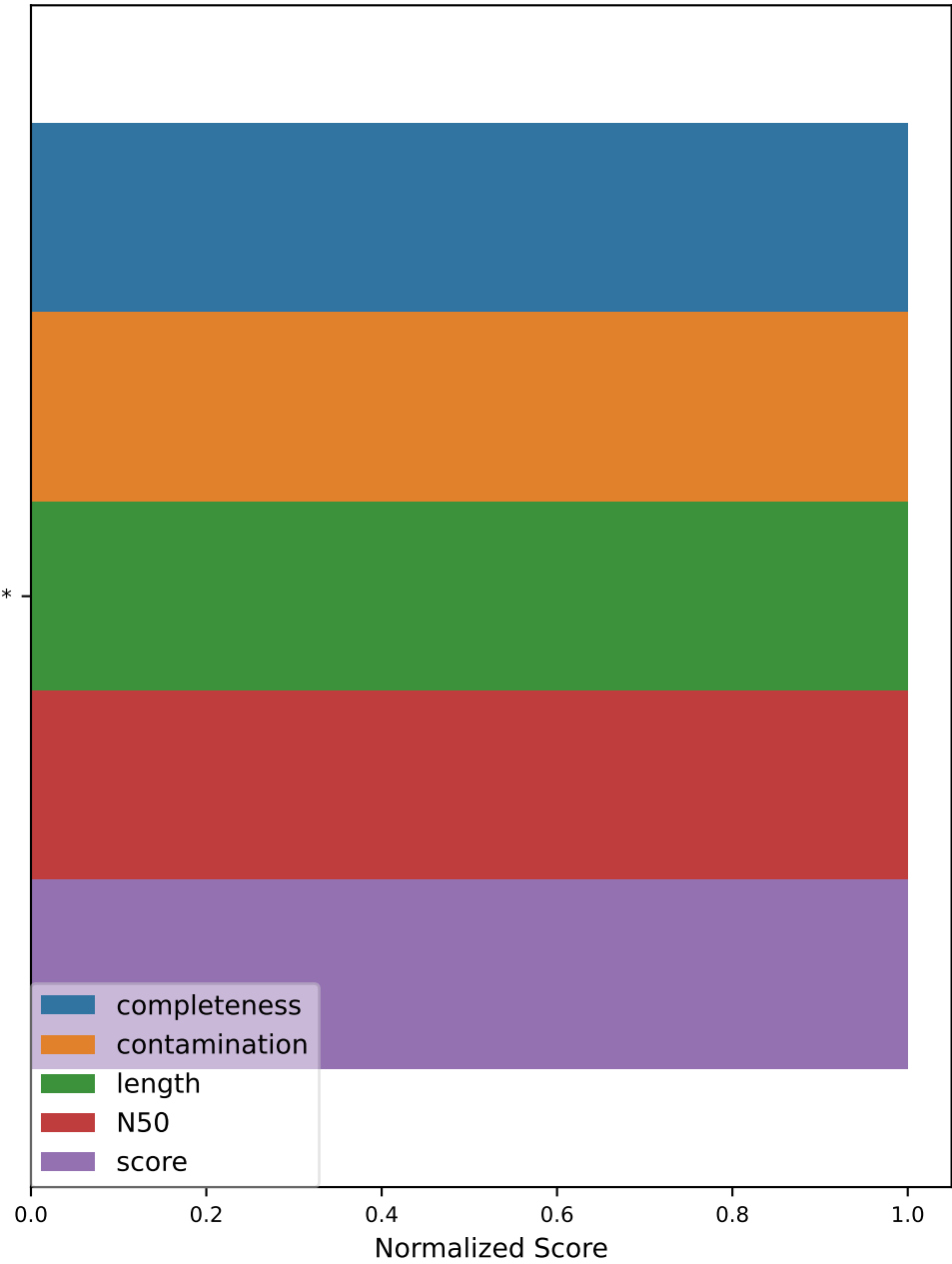
Scoring of cluster 60\_1



Scoring of cluster 61\_0

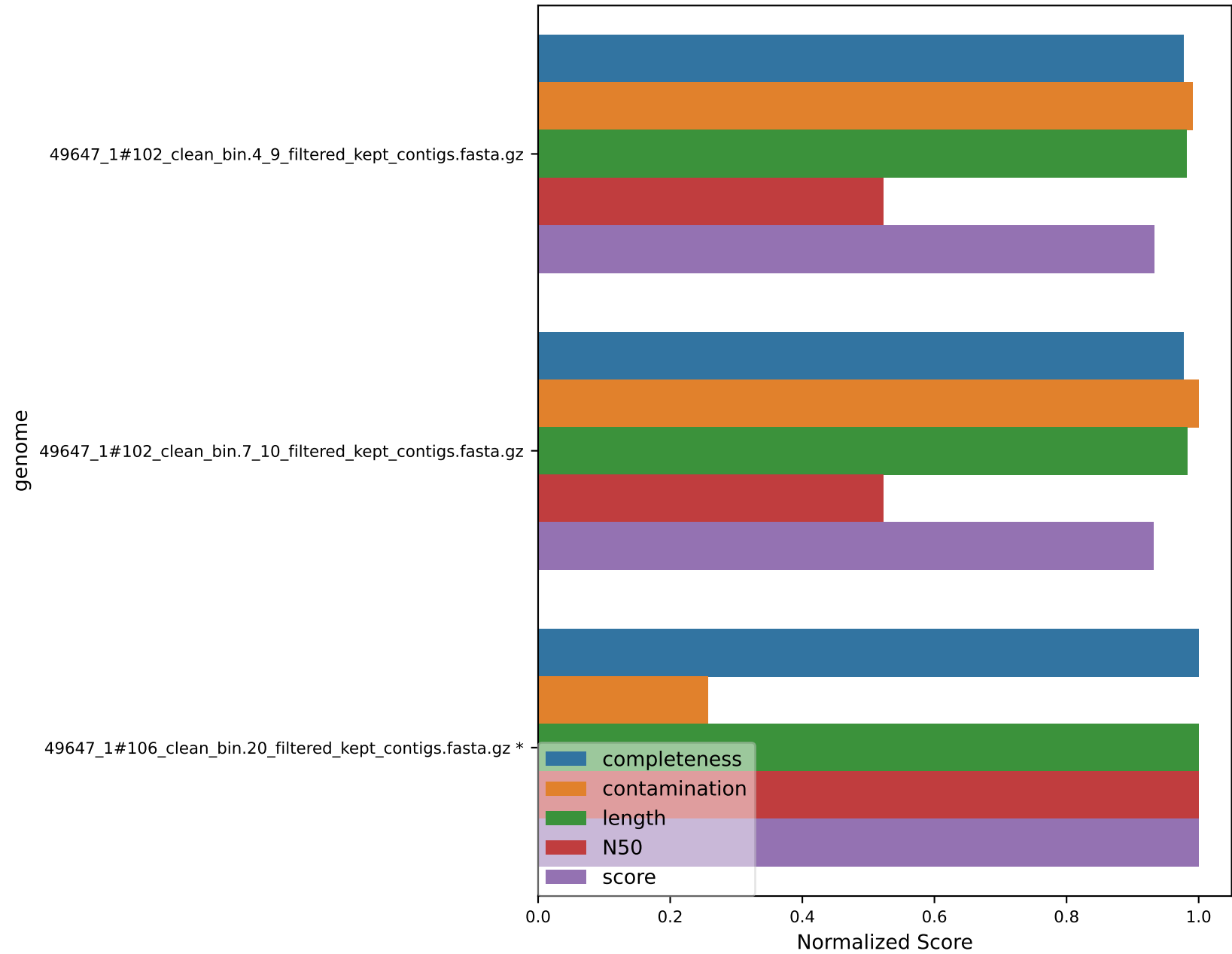
genome

49647\_1#2\_clean\_bin.25.orig\_filtered\_kept\_contigs.fasta.gz \*





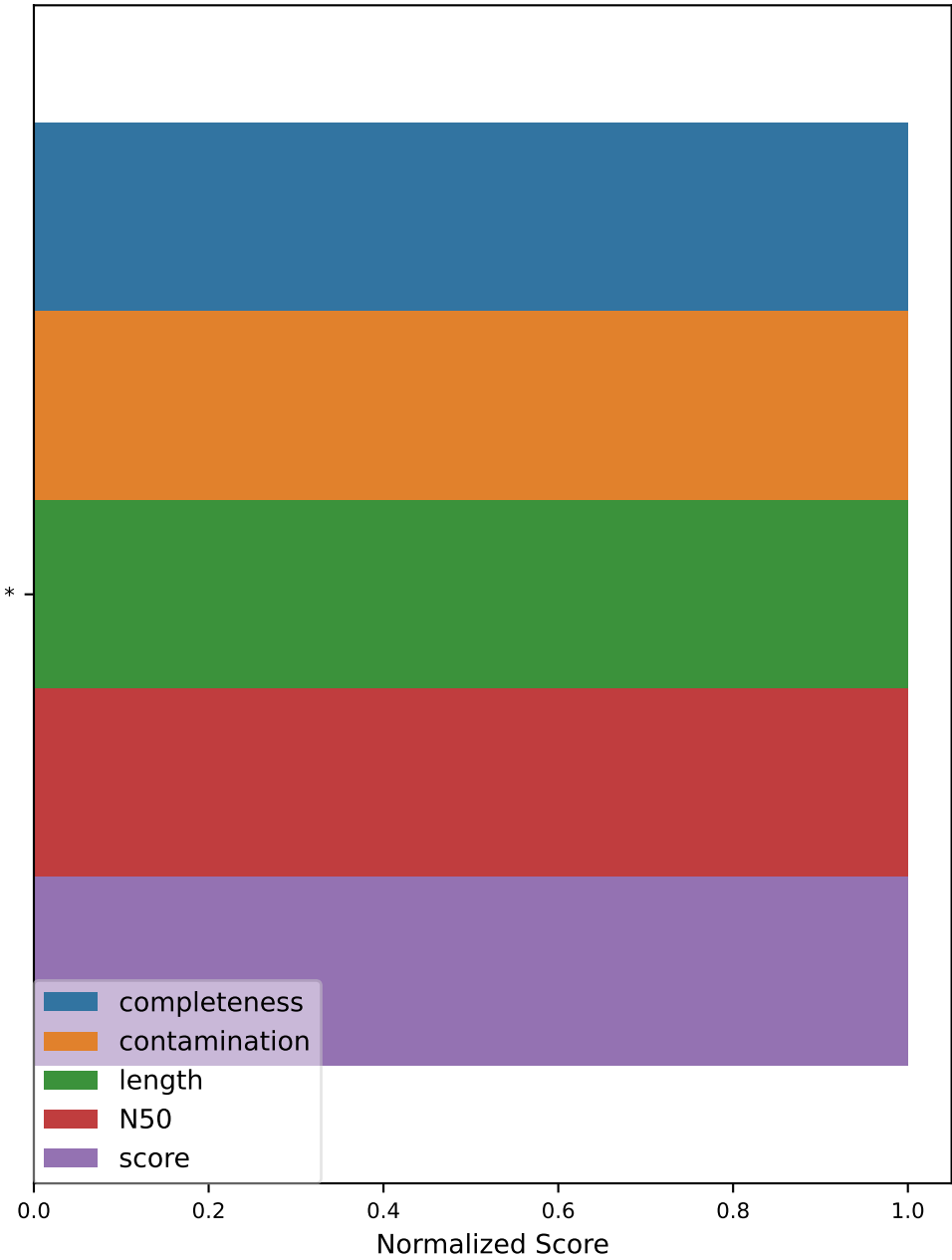
Scoring of cluster 62\_1



Scoring of cluster 62\_2

genome

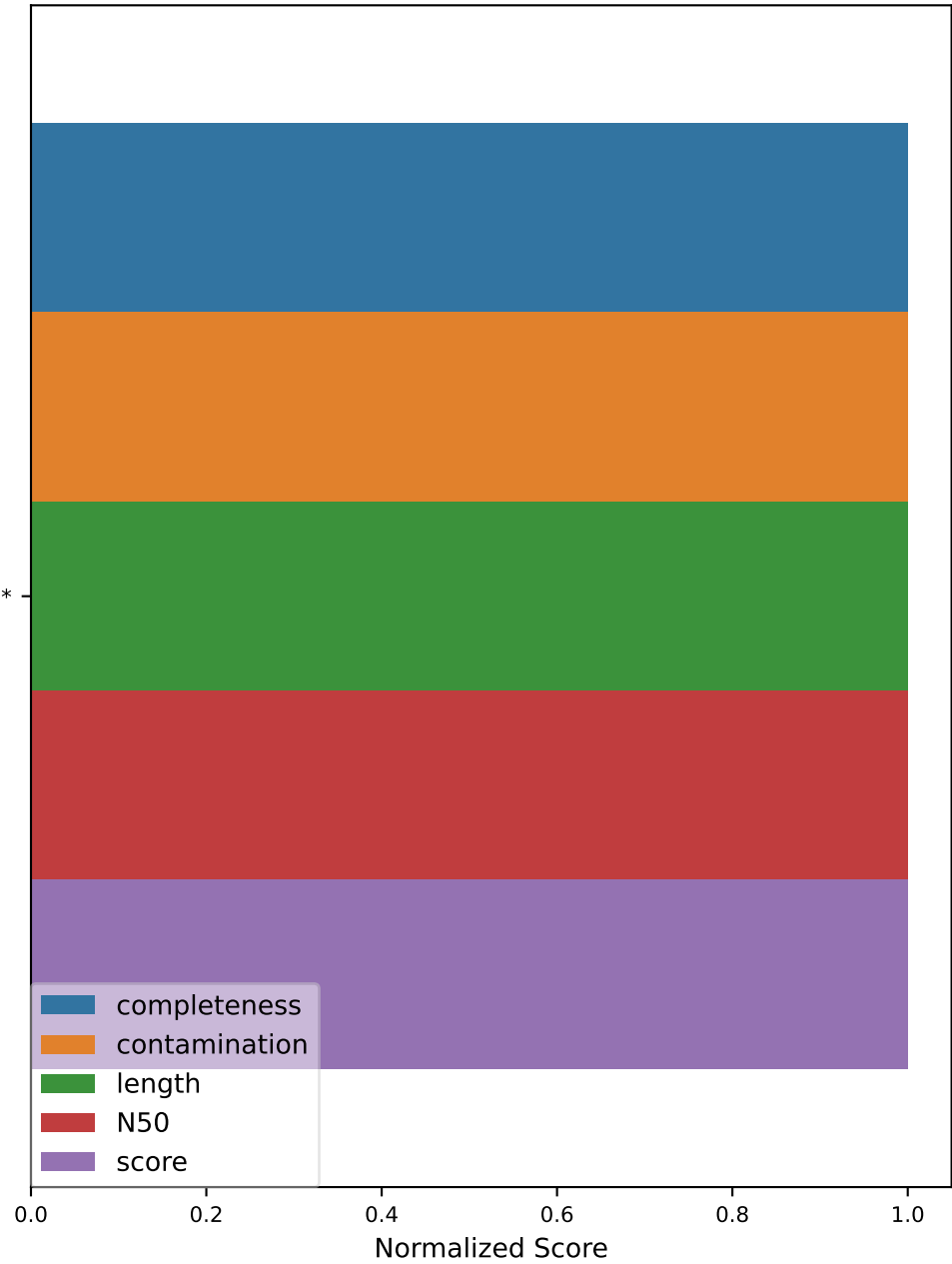
49647\_1#100\_clean\_bin.6\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 63\_0

genome

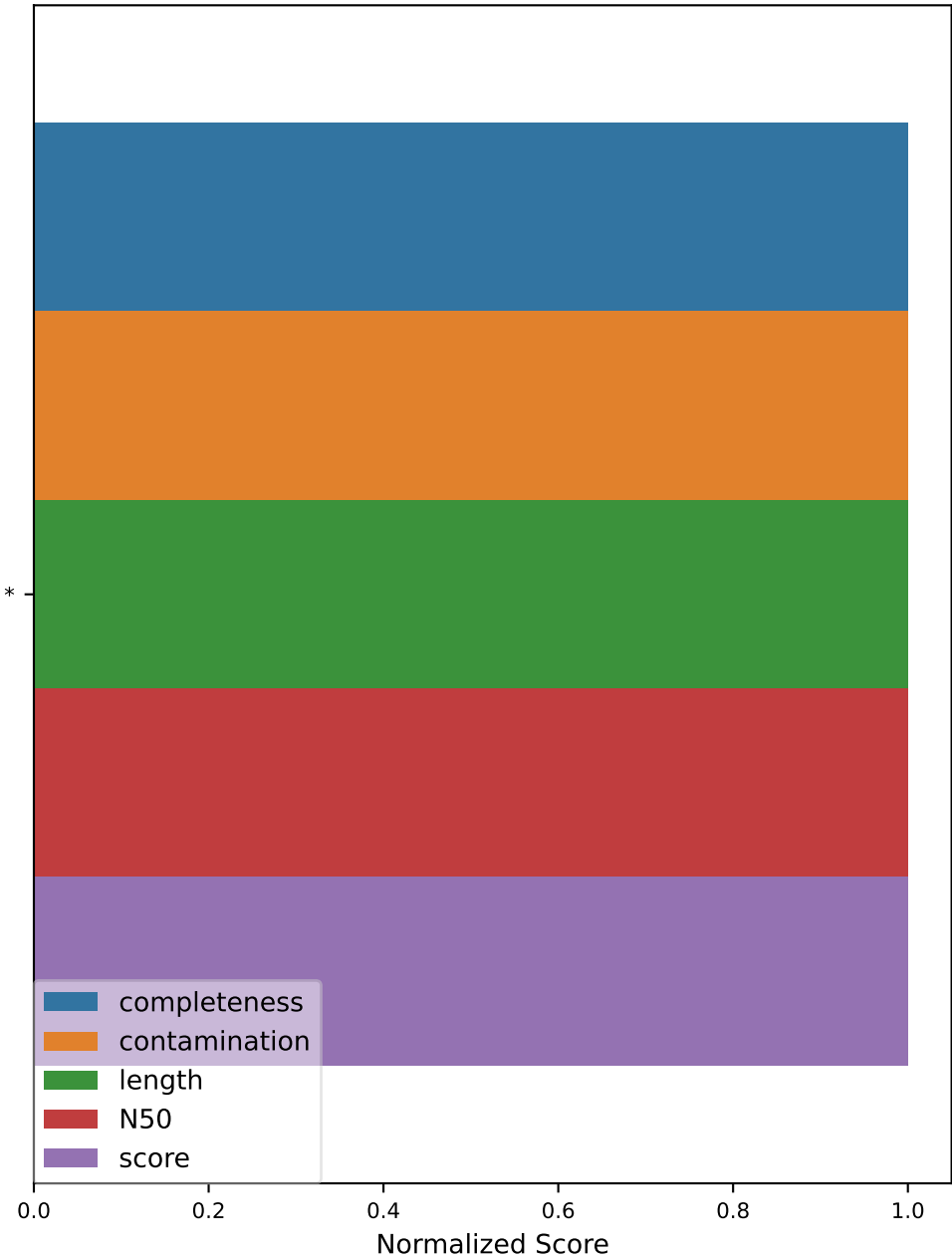
49647\_1#2\_clean\_bin.7.strict\_filtered\_kept\_contigs.fasta.gz \*



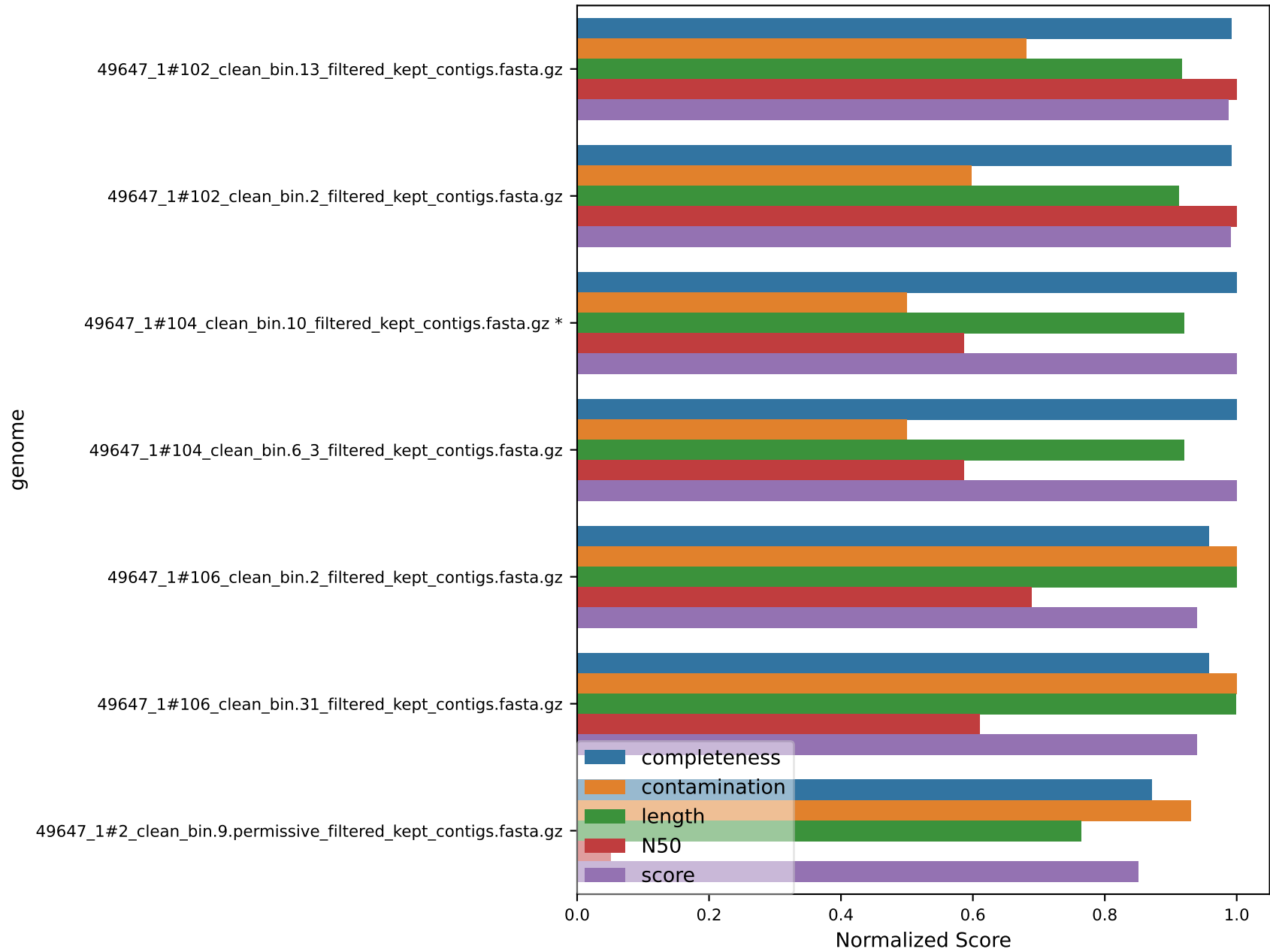
Scoring of cluster 64\_0

genome

49647\_1#106\_clean\_bin.22\_filtered\_kept\_contigs.fasta.gz \*



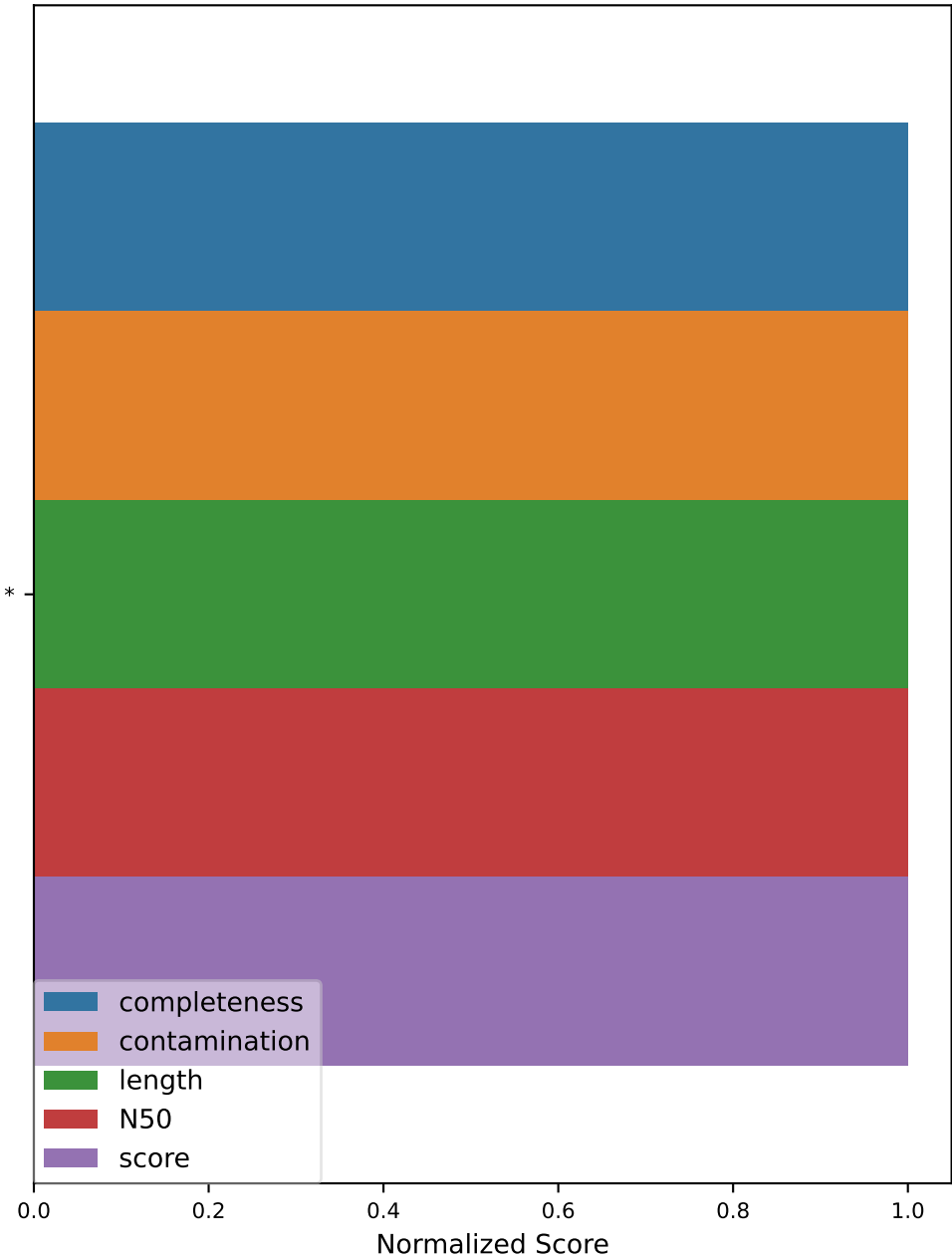
# Scoring of cluster 65\_1



Scoring of cluster 66\_0

genome

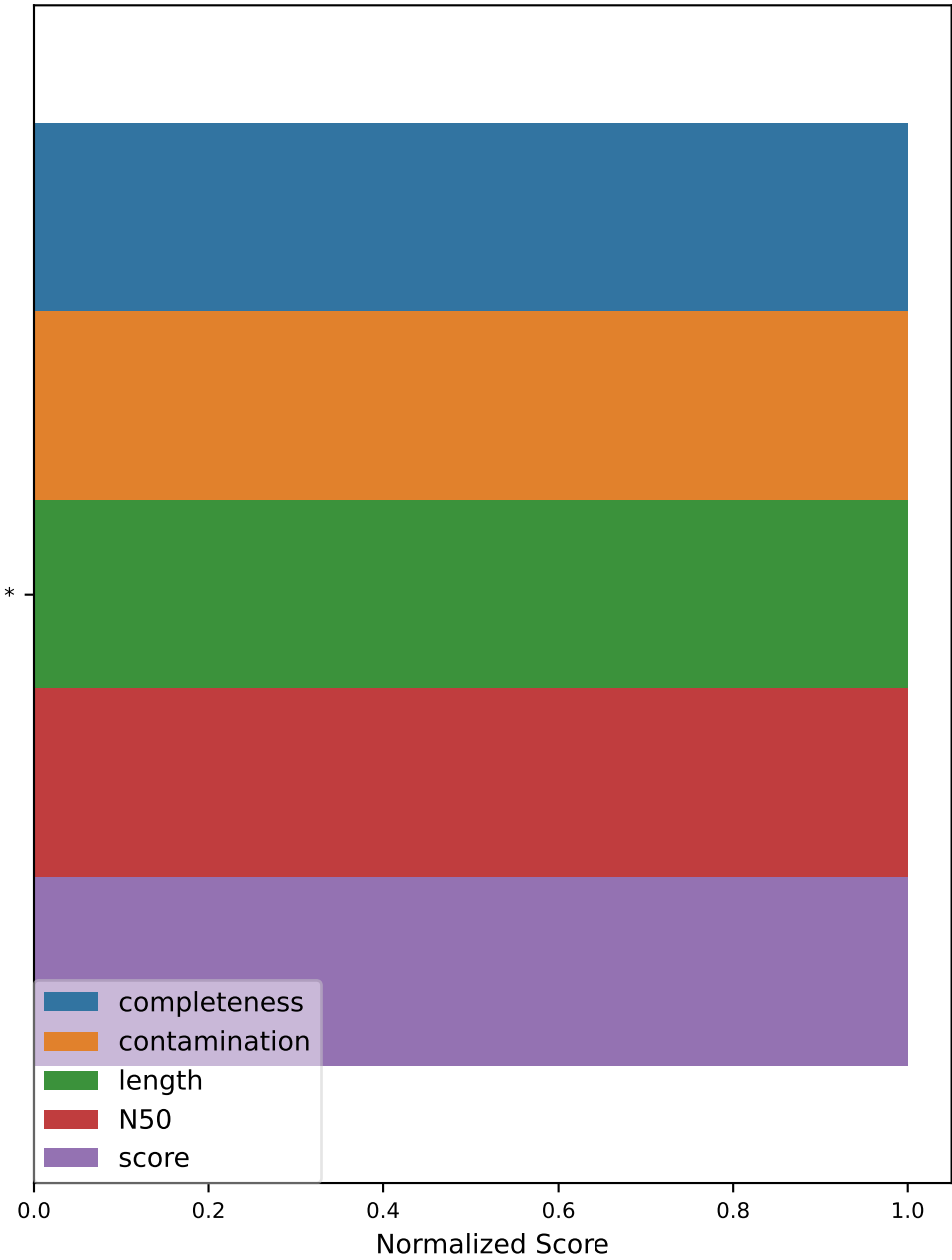
49647\_1#100\_clean\_bin.5\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 67\_0

genome

49647\_1#2\_clean\_bin.23.strict\_filtered\_kept\_contigs.fasta.gz \*



Scoring of cluster 68\_0

genome

49647\_1#2\_clean\_bin.29.strict\_filtered\_kept\_contigs.fasta.gz \*

