

# ERGA Assembly Report

v24.10.15

Tags: ERGA-BGE

TxID	3049732
ToLID	<b>qdStyilly1</b>
Species	<i>Stygiulus illyricus</i>
Class	Diplopoda
Order	Julida

Genome Traits	Expected	Observed
Haploid size (bp)	526,119,689	549,733,997
Haploid Number	12 (source: ancestor)	13
Ploidy	2 (source: ancestor)	2
Sample Sex	NA	NA

## EBP metrics summary and curation notes

Obtained EBP quality metric for pri: 6.7.Q52

The following metrics were automatically flagged as below EBP recommended standards or different from expected:

- . Observed Haploid Number is different from Expected
- . Not 90% of assembly in chromosomes for pri

## Curator notes

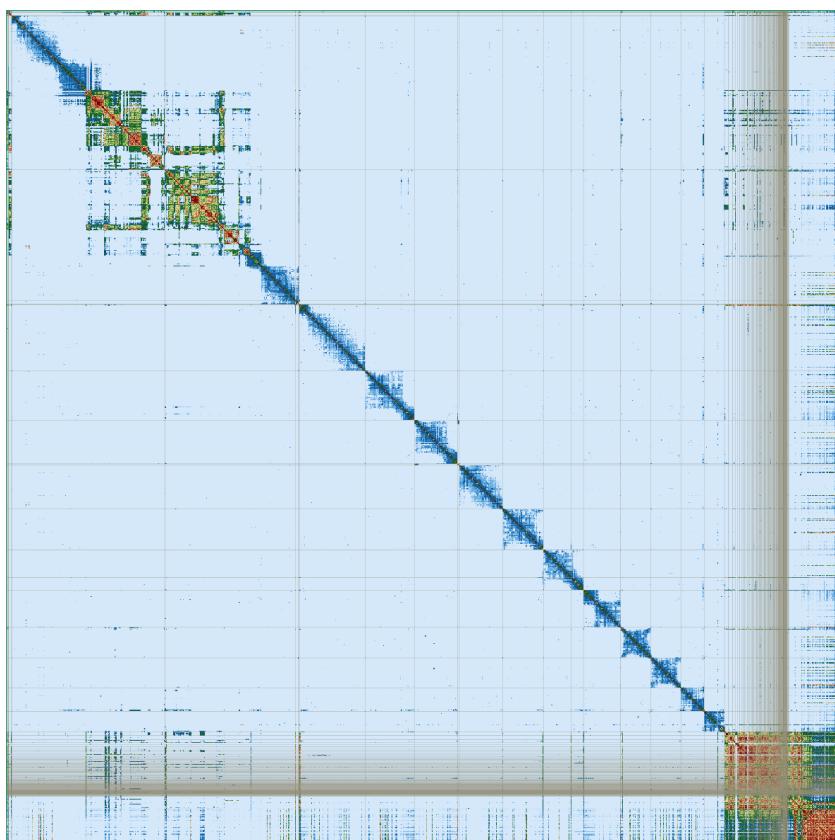
- . Interventions/Gb: None
- . Contamination notes: ""
- . Other observations: "Difficult to know if the first two scaffolds should be one or two chromosomes. Potentially the X and Y. Lots of repetitive sequences."

## Quality metrics table

Metrics	Pre-curation pri	Curated pri
Total bp	550,780,026	549,733,997
GC %	37.49	37.48
Gaps/Gbp	490.21	491.15
Total gap bp	54,000	54,000
Scaffolds	1,682	1,680
Scaffold N50	29,700,184	29,024,683
Scaffold L50	5	5
Scaffold L90	36	34
Contigs	1,952	1,950
Contig N50	2,585,720	2,585,720
Contig L50	60	60
Contig L90	235	234
QV	48.5773	52.7783
Kmer compl.	94.9653	87.5237
BUSCO sing.	97.1%	
BUSCO dupl.	0.6%	
BUSCO frag.	0.9%	
BUSCO miss.	1.4%	

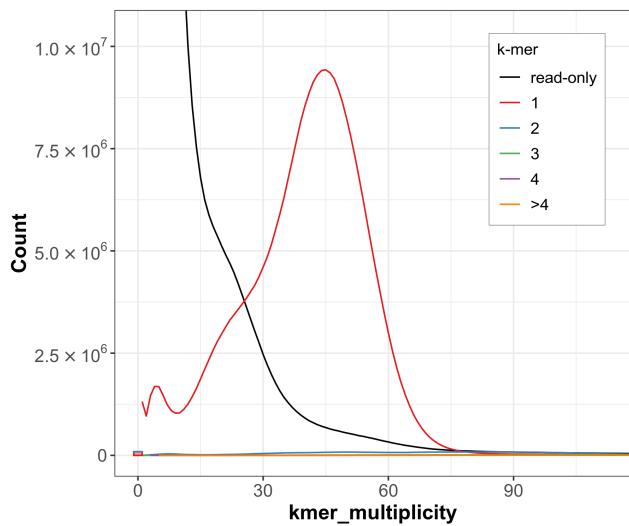
BUSCO: 6.0.0 (euk\_genome\_min, miniprot) / Lineage: arthropoda\_odb12 (genomes:76, BUSCOs:1667)

# HiC contact map of curated assembly

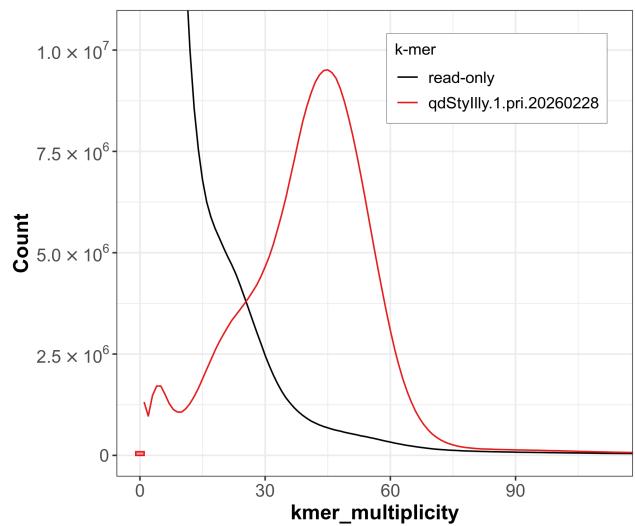


pri [LINK]

# K-mer spectra of curated assembly

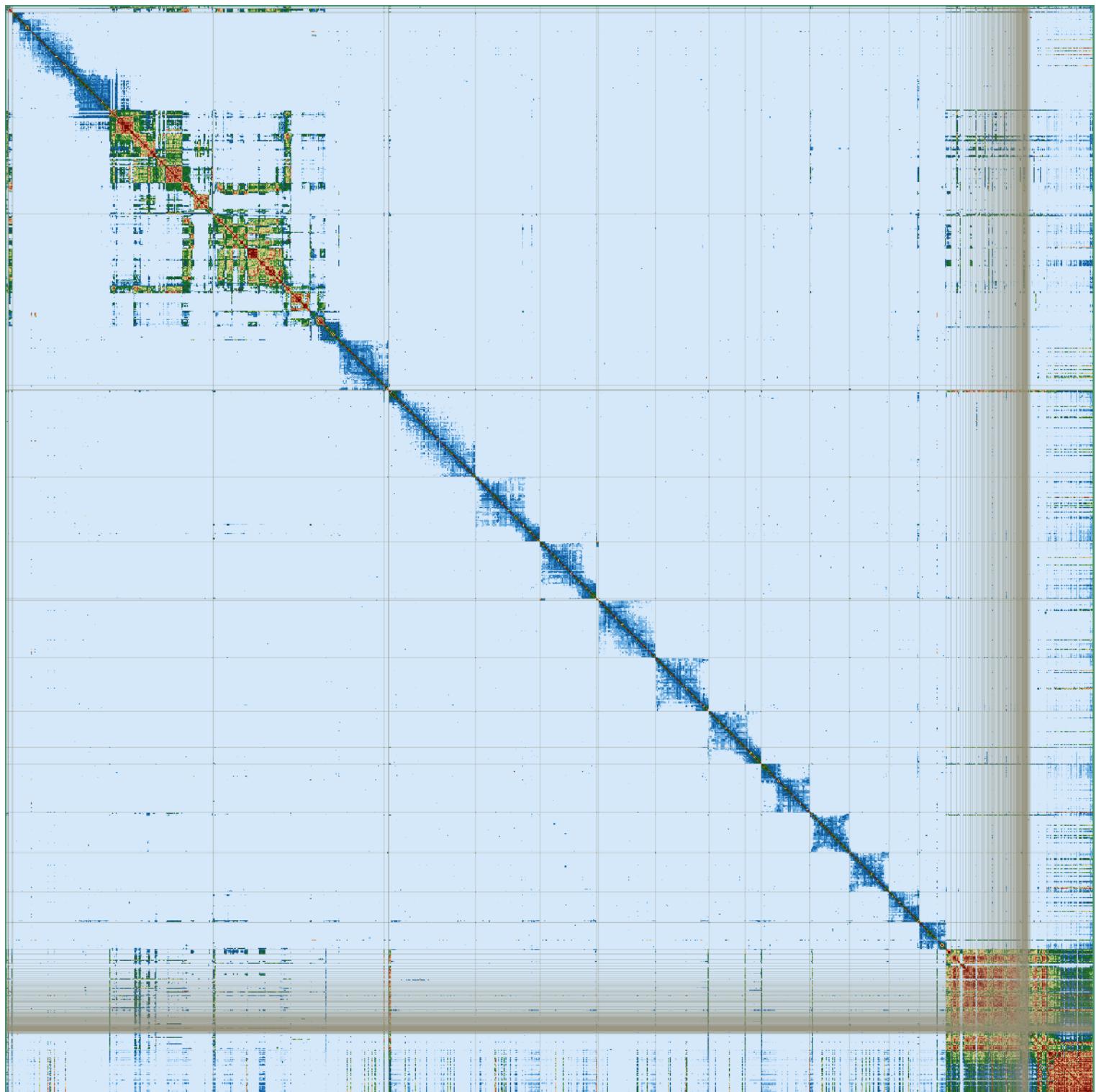


Distribution of k-mer counts per copy numbers found in asm



Distribution of k-mer counts coloured by their presence in reads/assemblies

## Post-curation contamination screening



**pri.** Bubble plot circles are scaled by sequence length, positioned by coverage and GC proportion, and coloured by taxonomy. Histograms show total assembly length distribution on each axis.

## Data profile

Data	ONT	HiC
Coverage	NA	NA

## Assembly pipeline

- **HiFiasm**
  - |\_ ver: 0.25.0
  - |\_ key param: NA
- **purge-dups**
  - |\_ ver: 1.2.6
  - |\_ key param: NA
- **yahs**
  - |\_ ver: 1.1a
  - |\_ key param: NA
- **fcs-gx**
  - |\_ ver: 0.5.0
  - |\_ key param: NA

## Curation pipeline

- **PretextView**
  - |\_ ver: 1.0.0
  - |\_ key param: NA

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