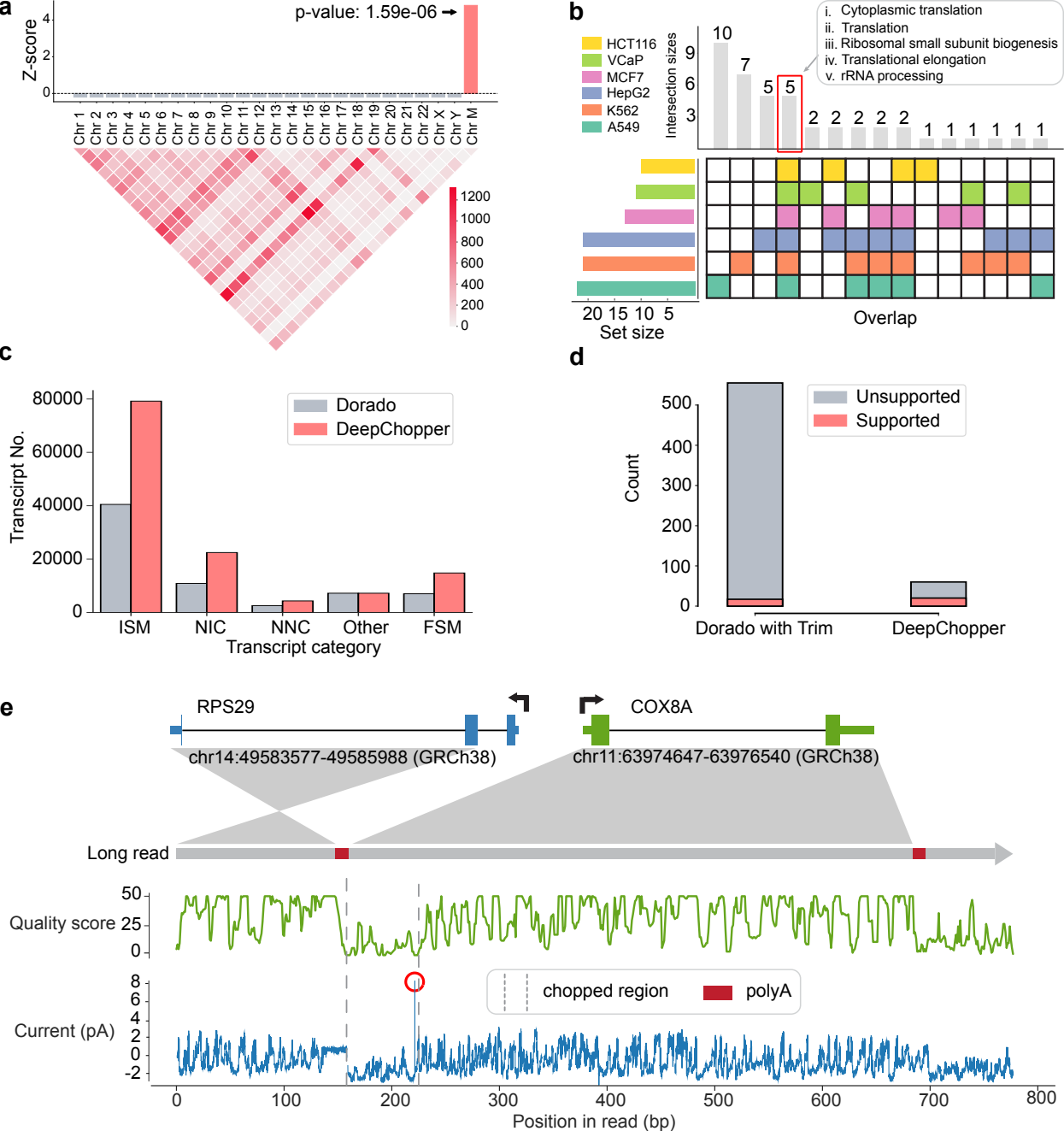
a
p-value: 1.59e-06
Z-score
Chr 1
Chr 2
Chr 3
Chr 4
Chr 5
Chr 6
Chr 7
Chr 8
Chr 9
Chr 10
Chr 11
Chr 12
Chr 13
Chr 14
Chr 15
Chr 16
Chr 17
Chr 18
Chr 19
Chr 20
Chr 21
Chr 22
Chr X
Chr Y
Chr M
b
HCT116
VCaP
MCF7
HepG2
K562
A549
Intersection sizes
i. Cytoplasmic translation
ii. Translation
iii. Ribosomal small subunit biogenesis
iv. Translational elongation
v. rRNA processing
Set size
Overlap
c
Transcirpt No.
Dorado
DeepChopper
ISM
NIC
NNC
Other
FSM
Transcript category
d
Count
Unsupported
Supported
Dorado with Trim
DeepChopper
e
RPS29
chr14:49583577-49585988 (GRCh38)
COX8A
chr11:63974647-63976540 (GRCh38)
Long read
Quality score
Current (pA)
chopped region
polyA
Position in read (bp)