Challenge 3 Evaluation Predictions Entry Entry Entry submission 4 submission 7 submission 1 **Ground truth** Entry Entry submission 5 submission 2 Available before submission Entry Entry **Entry** SIRV Set 3 (manatee) submission 6 submission 3 submission N SIRV Set 4 (mouse) **SQANTI3** Evaluation **Comparative assessment** Similar comparisons to Challenge 1 but Transcript length only with SIRVs Illumina support high mapping rate to genome high number of complete BUSCO genes Compare Compare sequencing platform pipelines Select candidate targets for further experimental validation