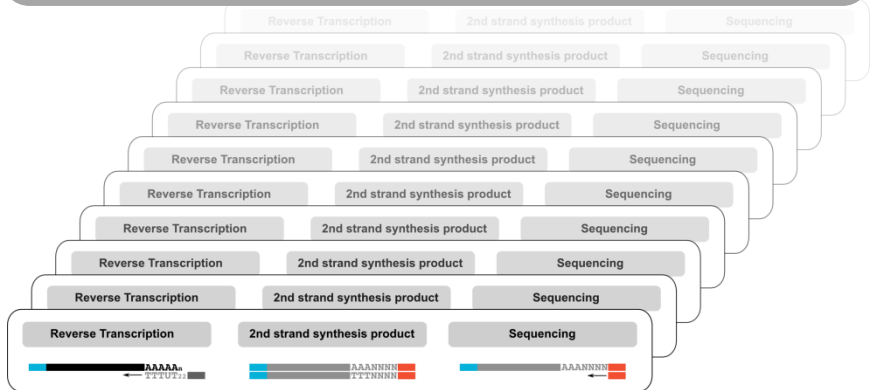


**Different 3' end sequencing libraries...**



### 3' end sequencing libraries

human: 78 samples (7 protocols)

mouse: 110 samples (8 protocols)

human: 78 samples (7 protocols)

mouse: 110 samples (8 protocols)



The image shows a screenshot of the PolyASite website. The top section has a blue background with the text "PolyASite" in a large, stylized font on the left, and "Welcome to PolyASite" followed by "Repository for 3' end sequencing data" on the right. Below this, a light blue banner contains the text "Check out our 'About' section to easily figure out how to get the data you want." The bottom section features a navigation bar with five icons and labels: a home icon labeled "Home", a flask icon labeled "Browse by experiment", a DNA sequence icon labeled "3' end sequencing techniques", a globe icon labeled "Poly(A) sites atlas", and a speech bubble icon labeled "About". At the very bottom, a grey box contains the text "Number of poly(A) sites per species:" followed by "Mouse: 183,225" and "Human: 392,912".

Repository for 3' end sequencing data

Check out our "About" section to easily figure out how to get the data you want.

[Home](#)

Number of poly(A) sites per species:

Mouse: 183,225

Human: 392,912

### Uniform processing pipeline

- protocol-specific library preprocessing
- remove protocol specific artifacts

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