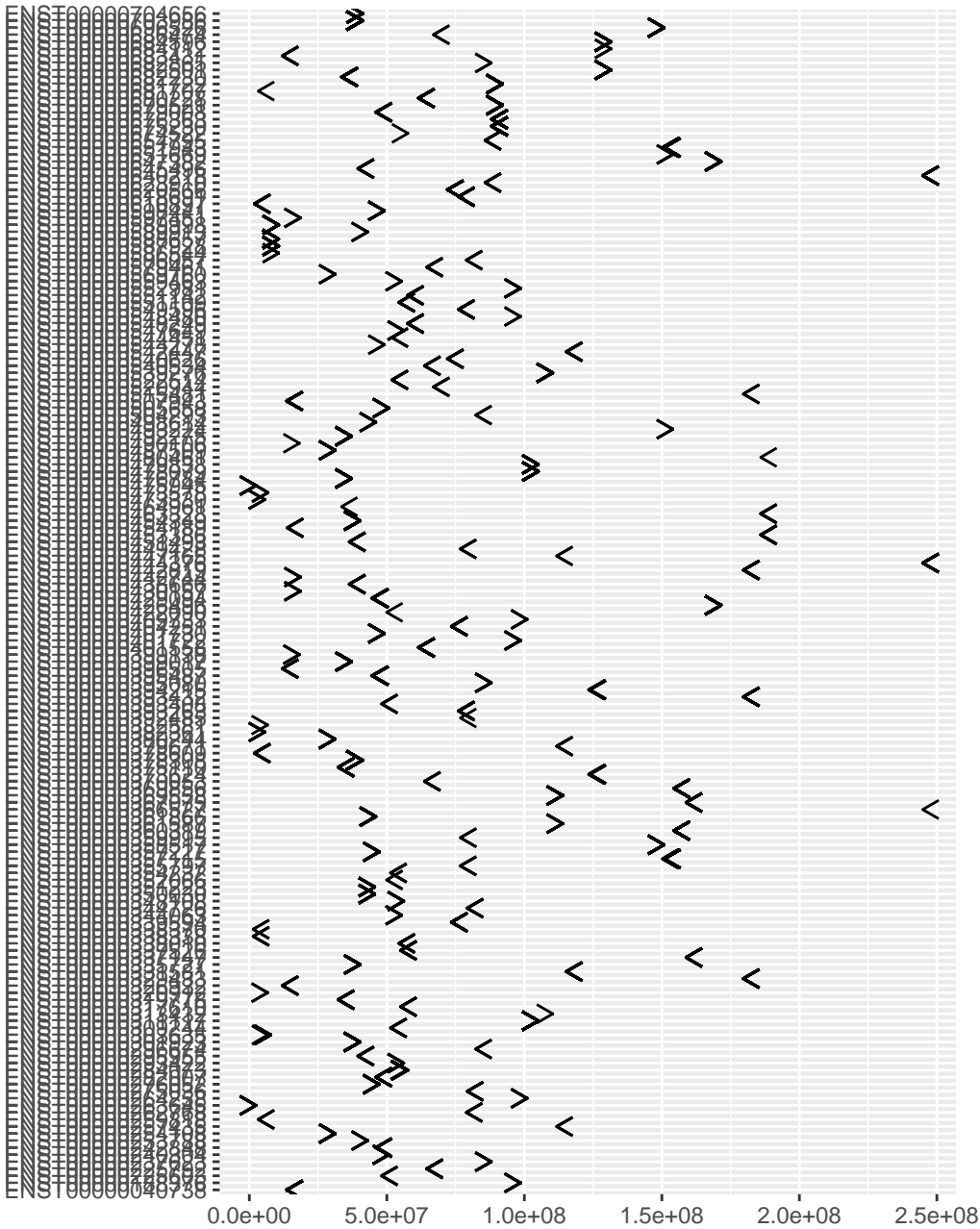


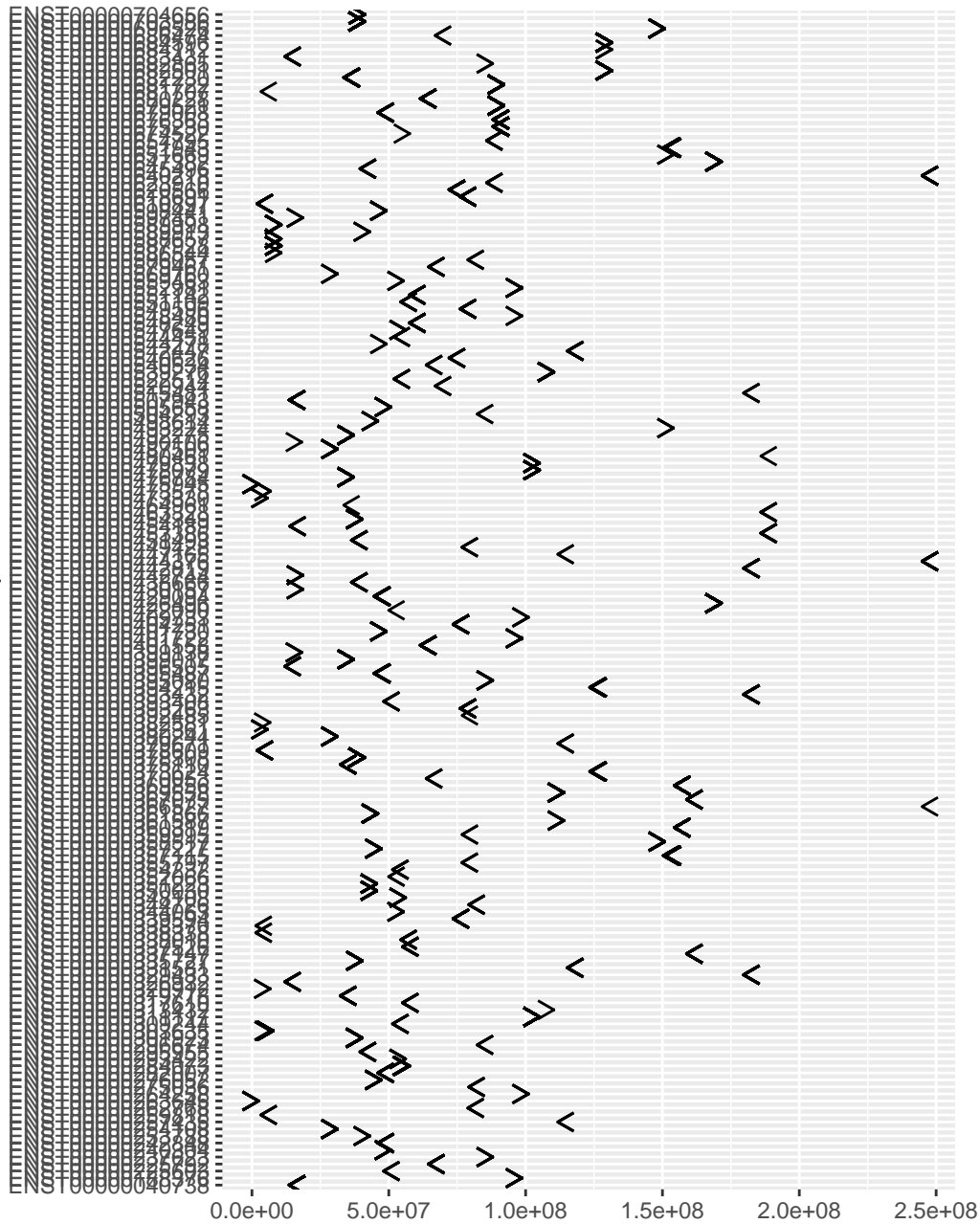
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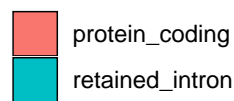


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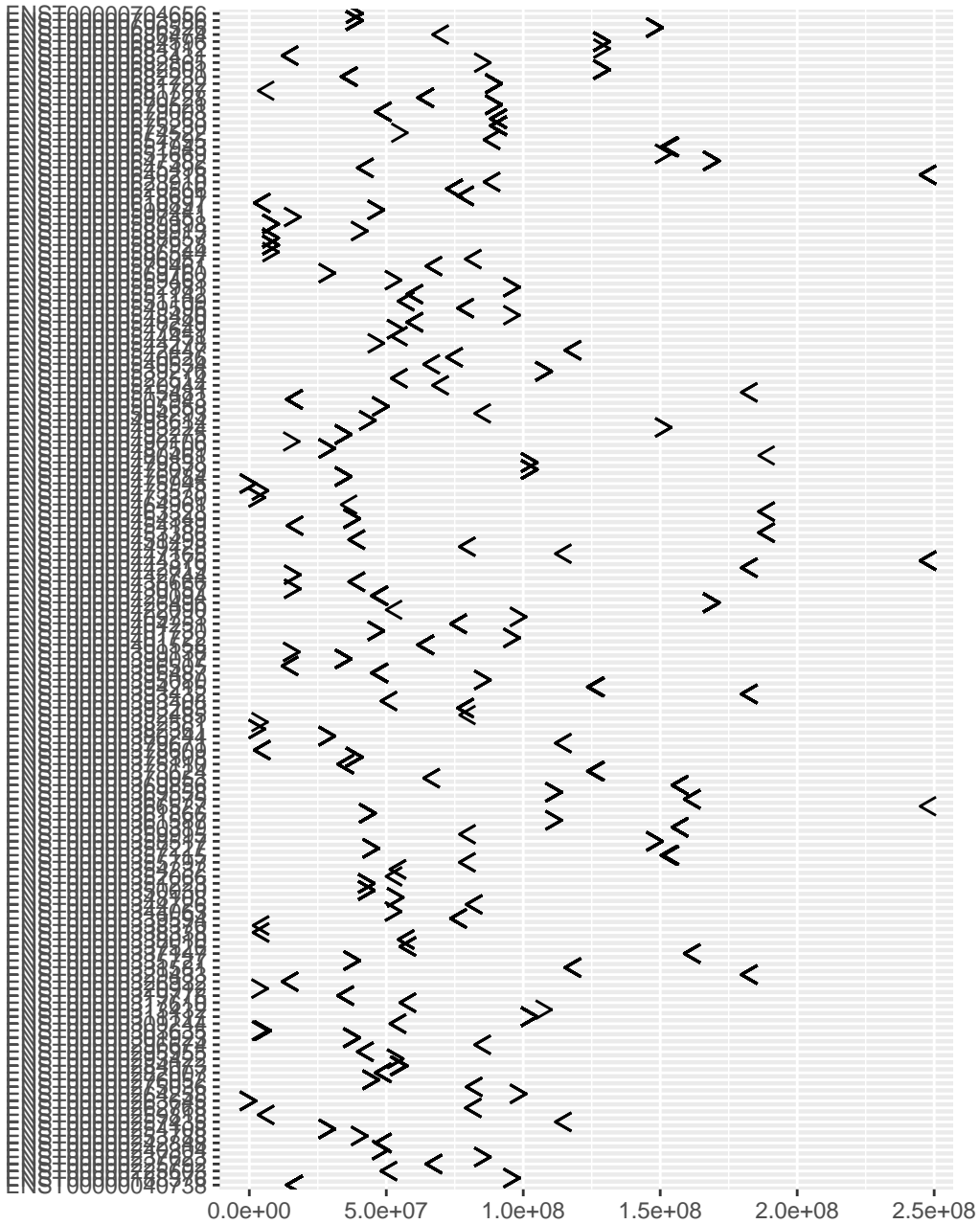


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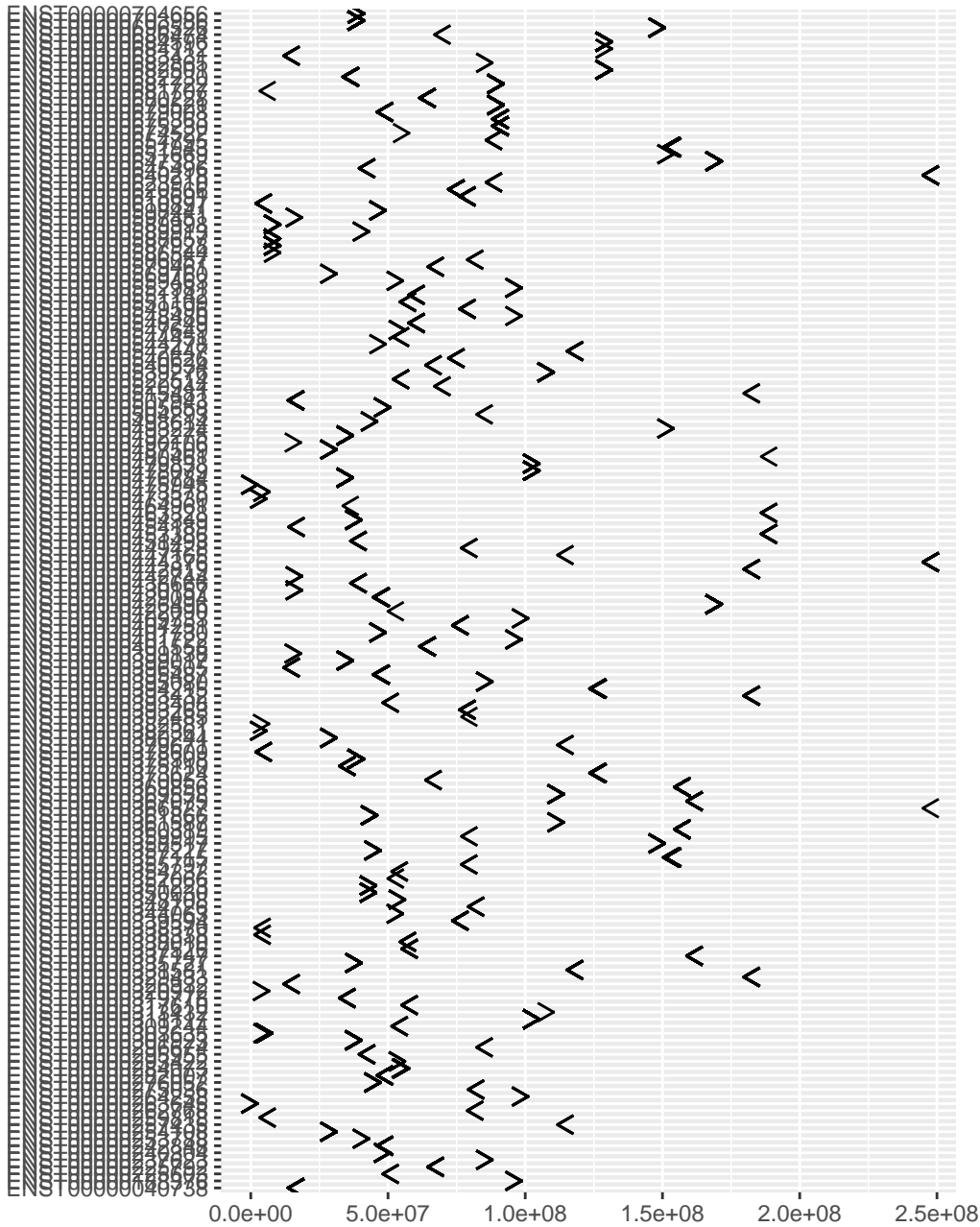
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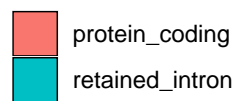
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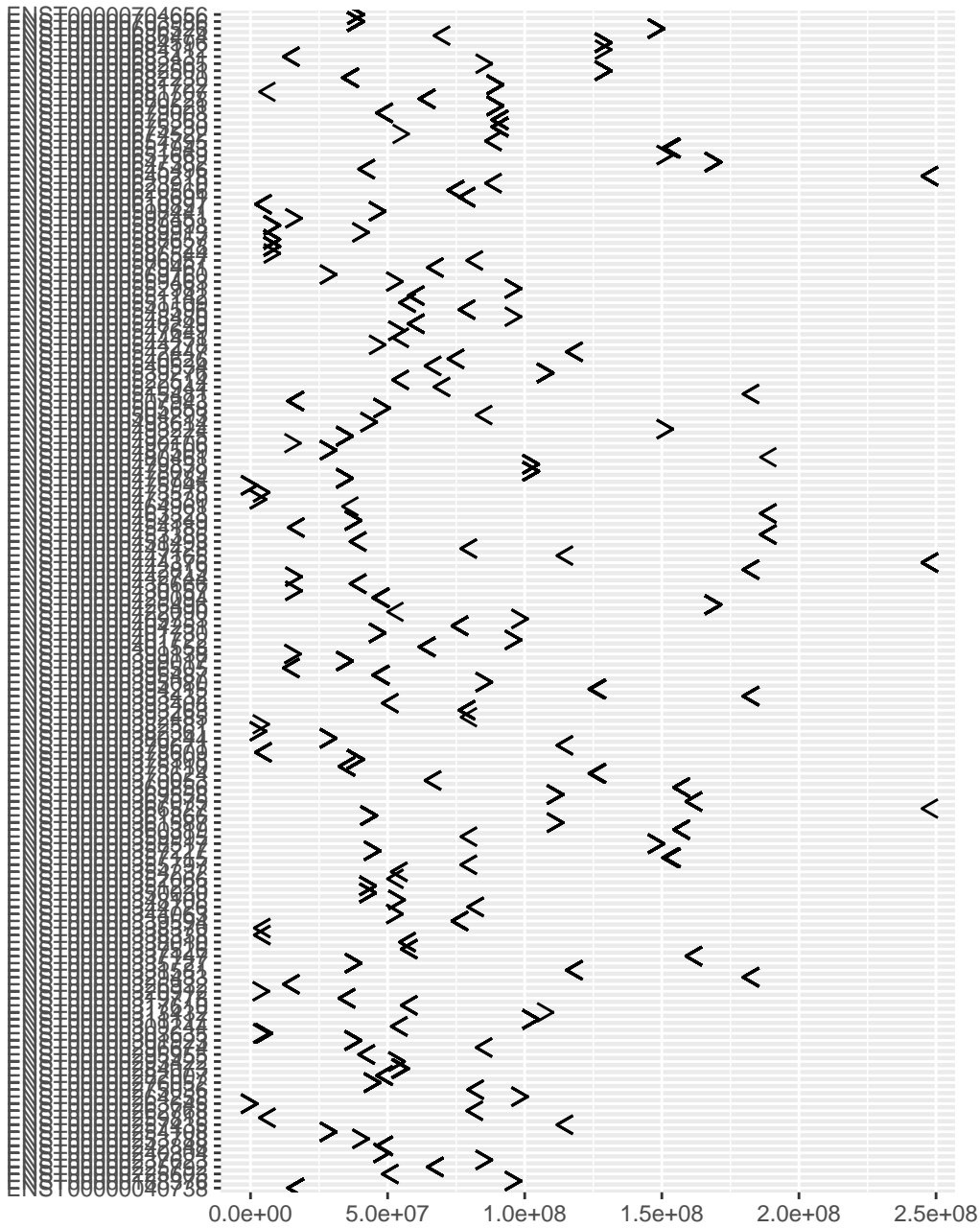


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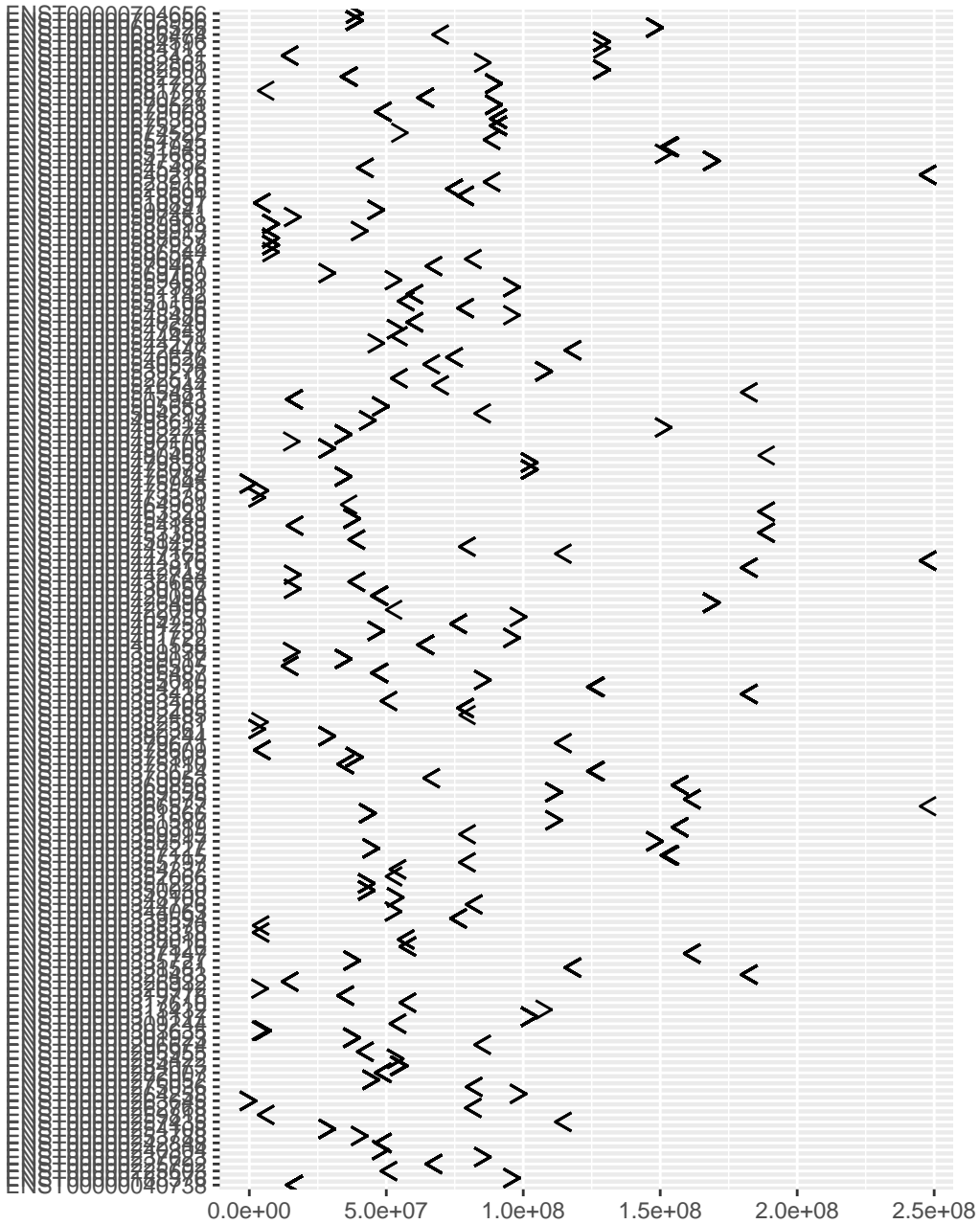
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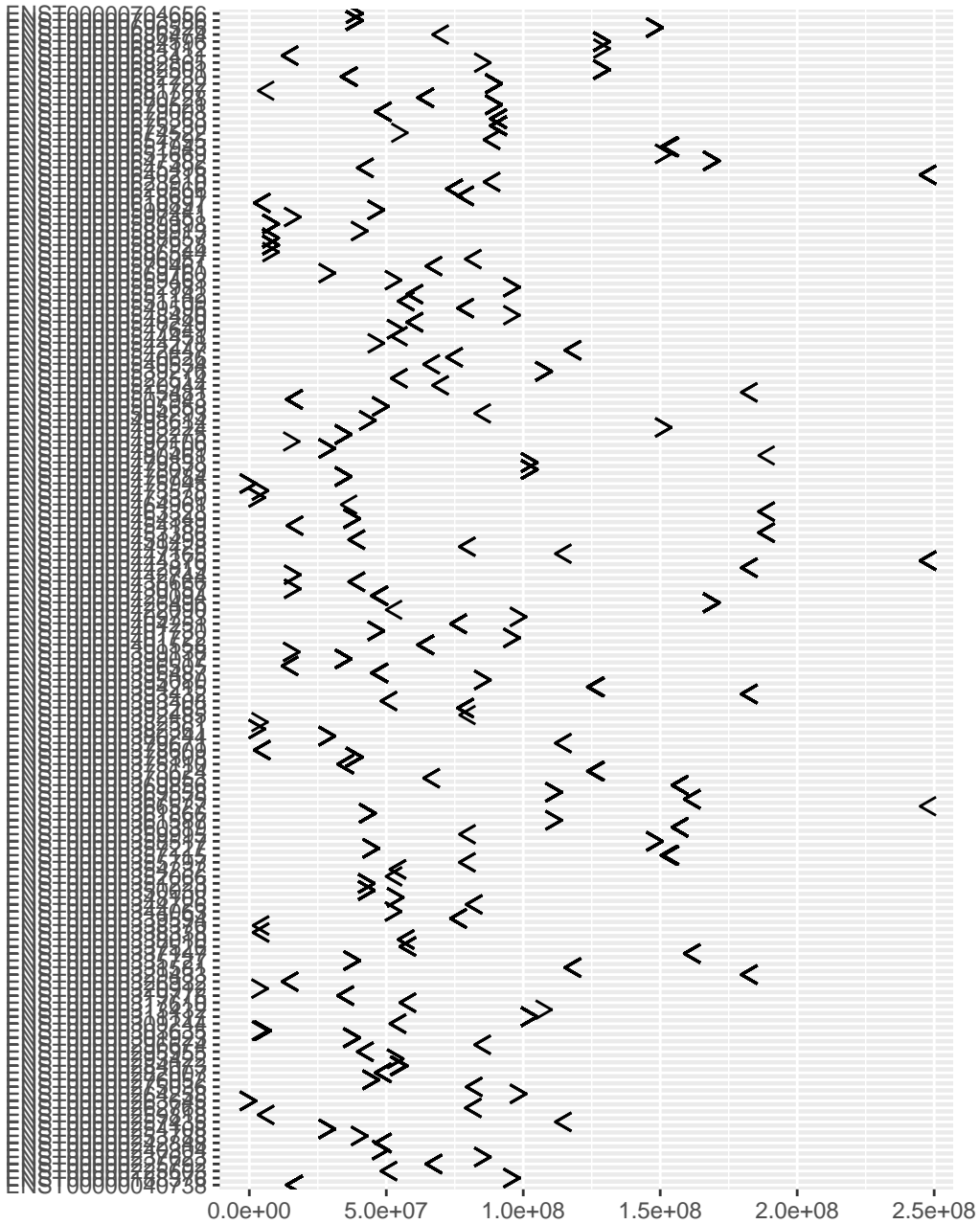
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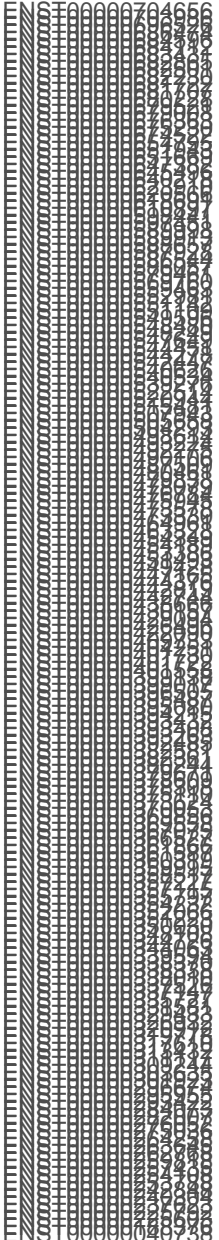
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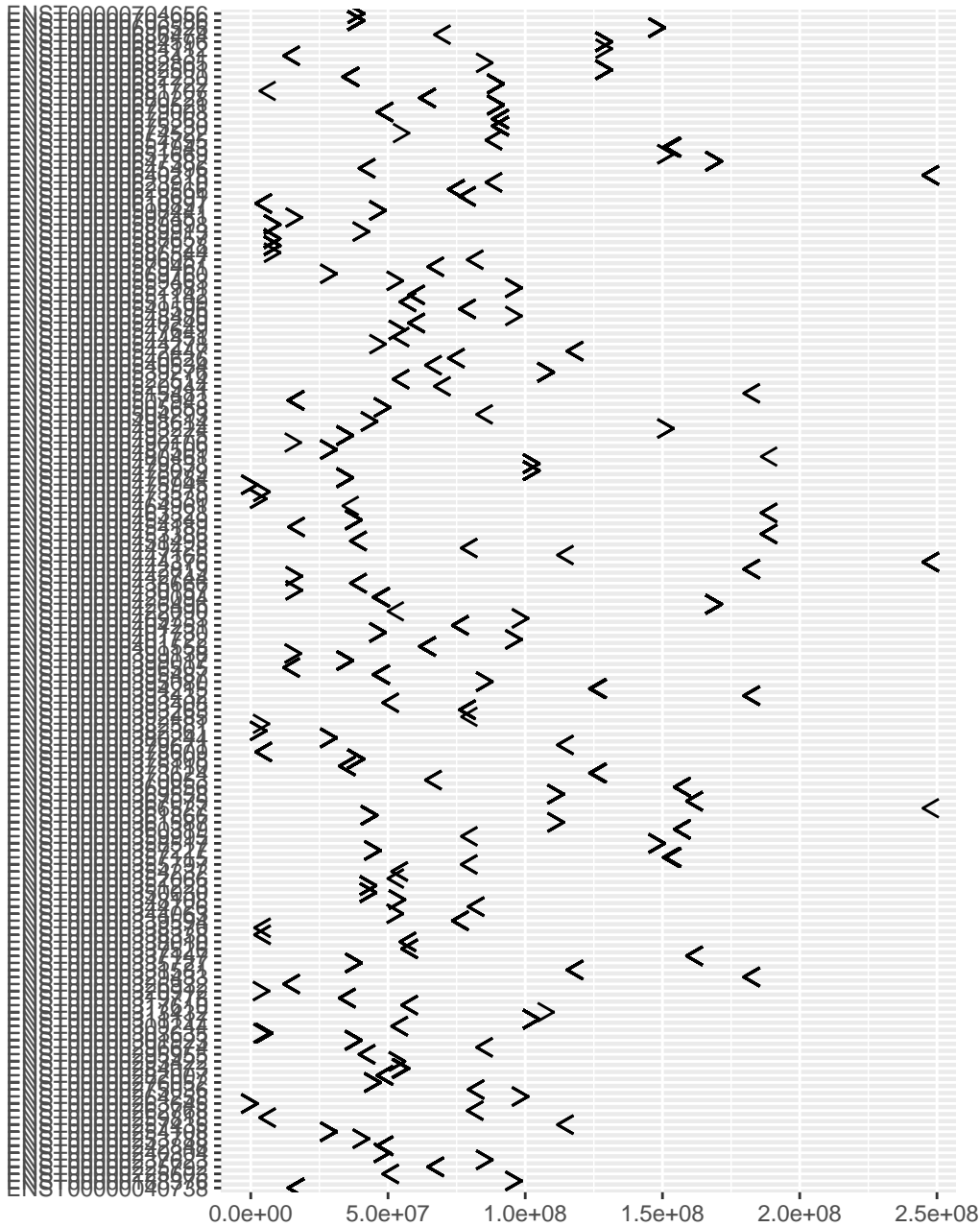


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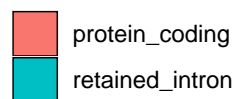


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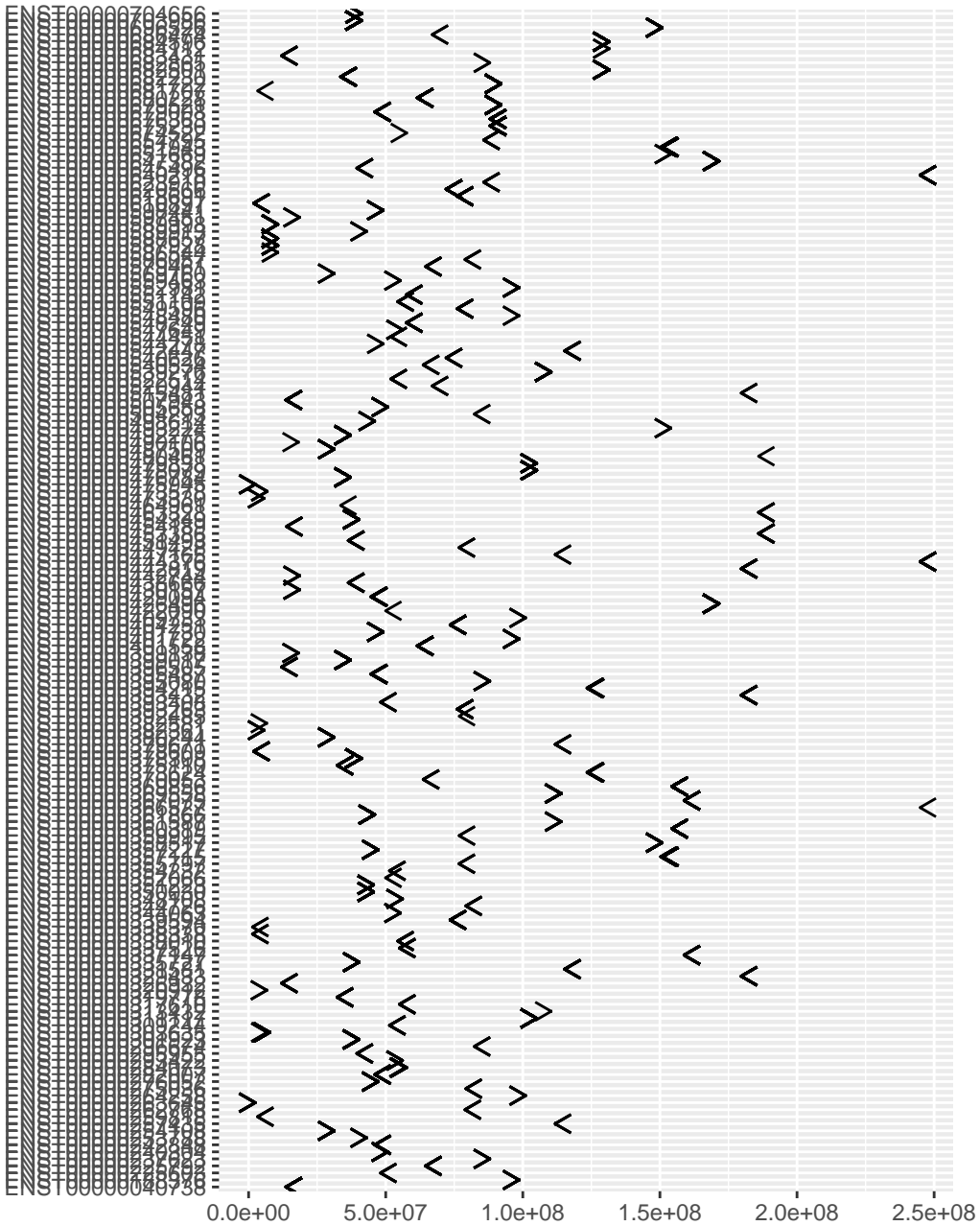


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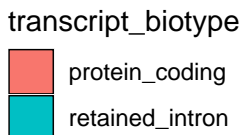
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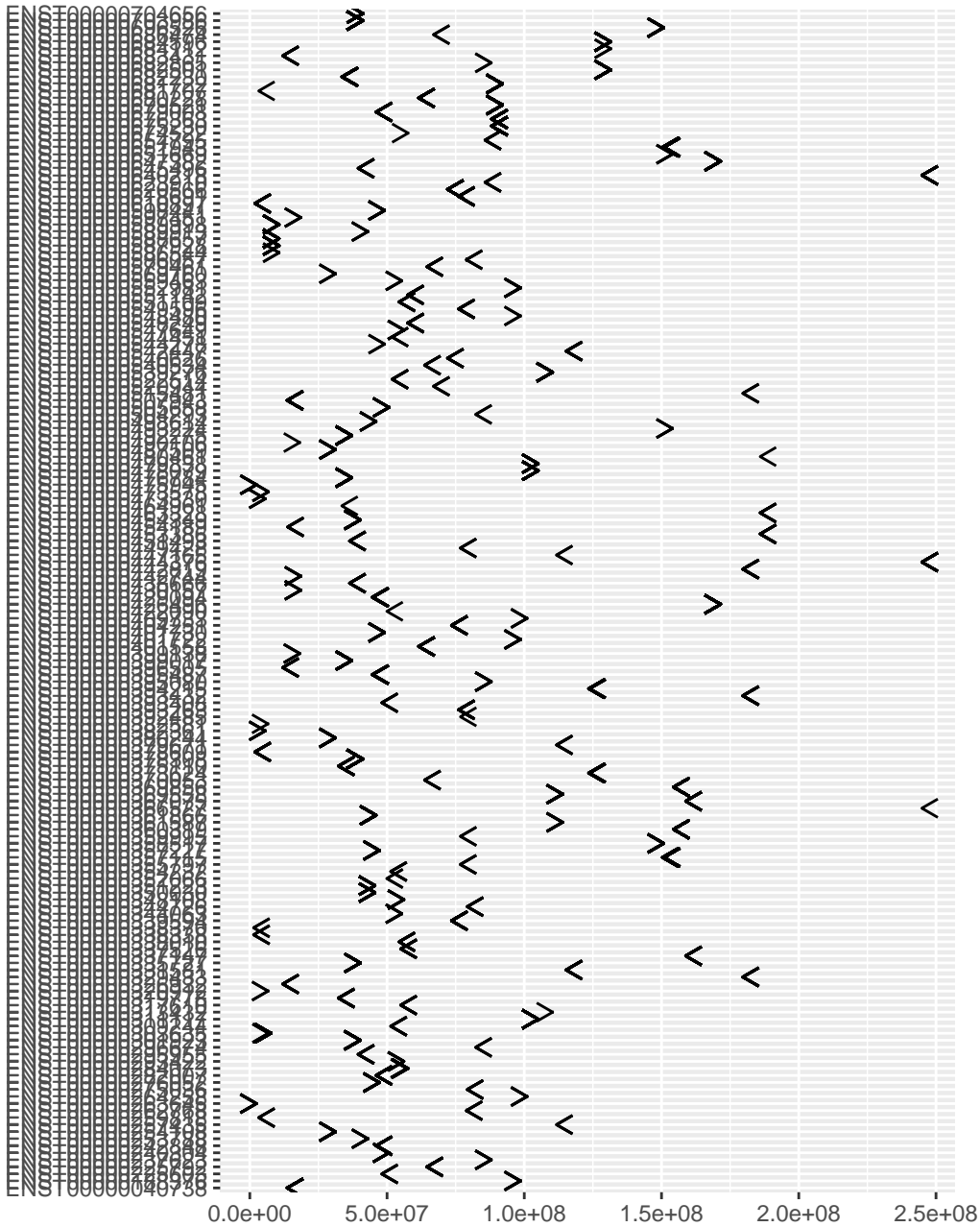
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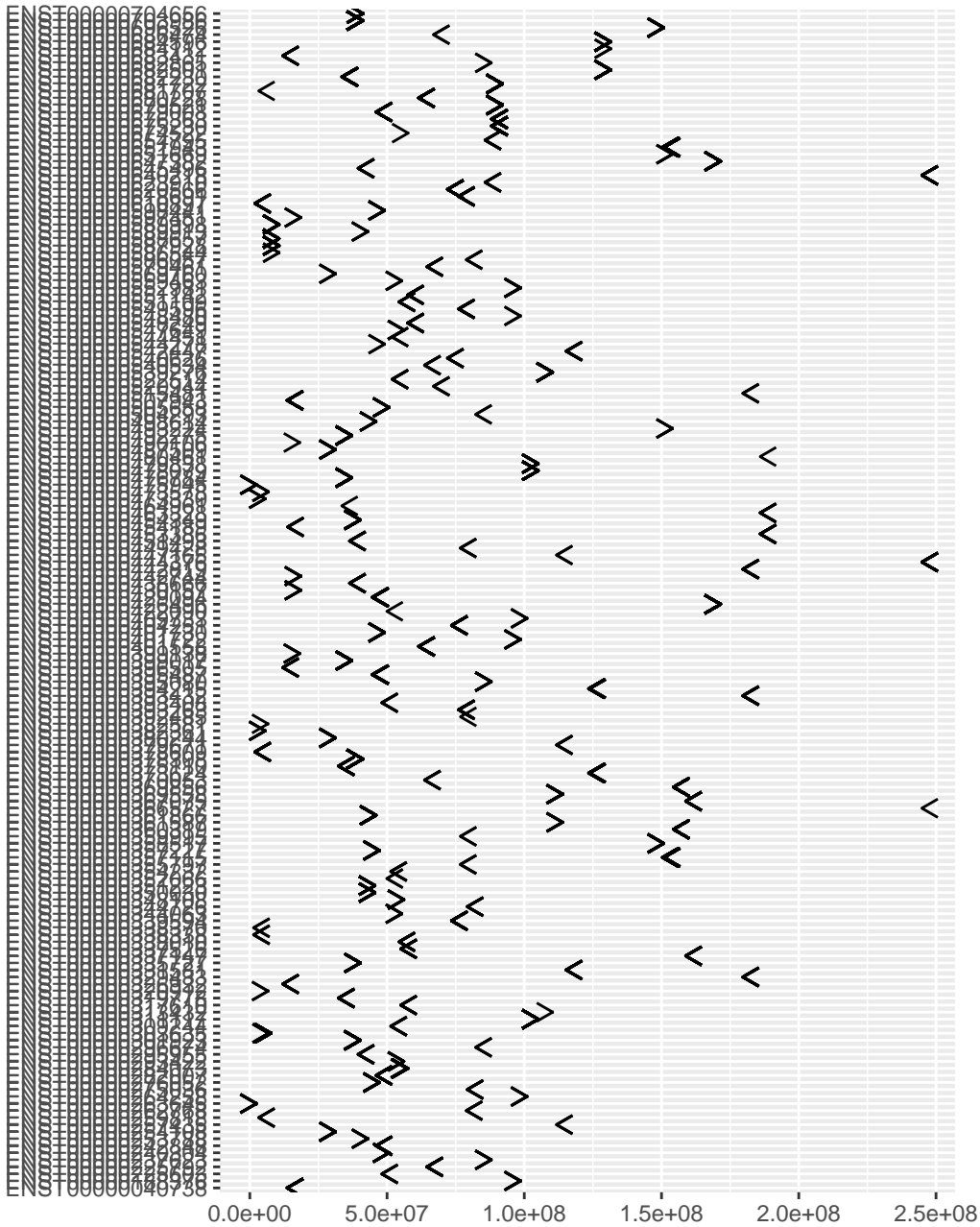
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protein_coding

CD81

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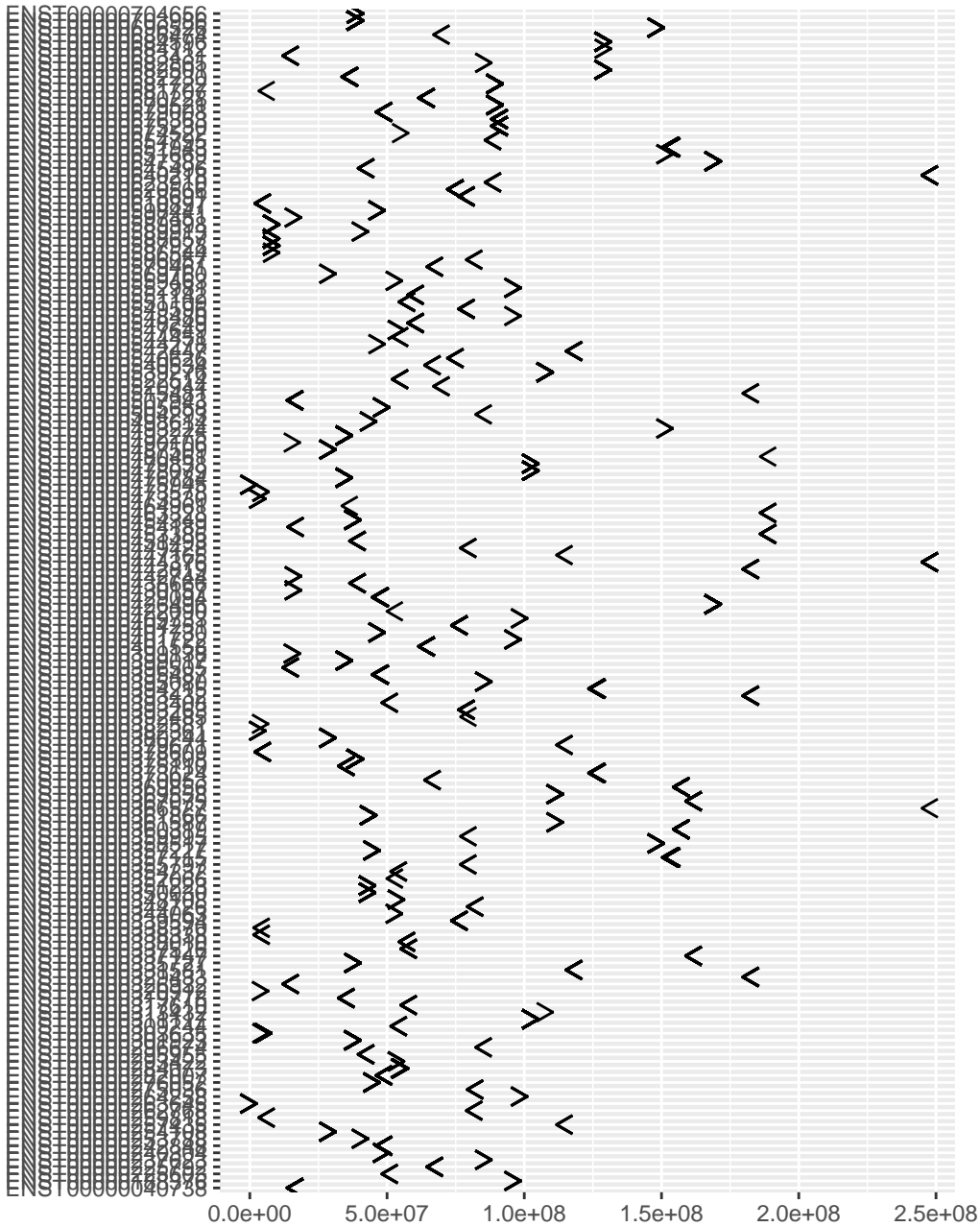
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RPL31

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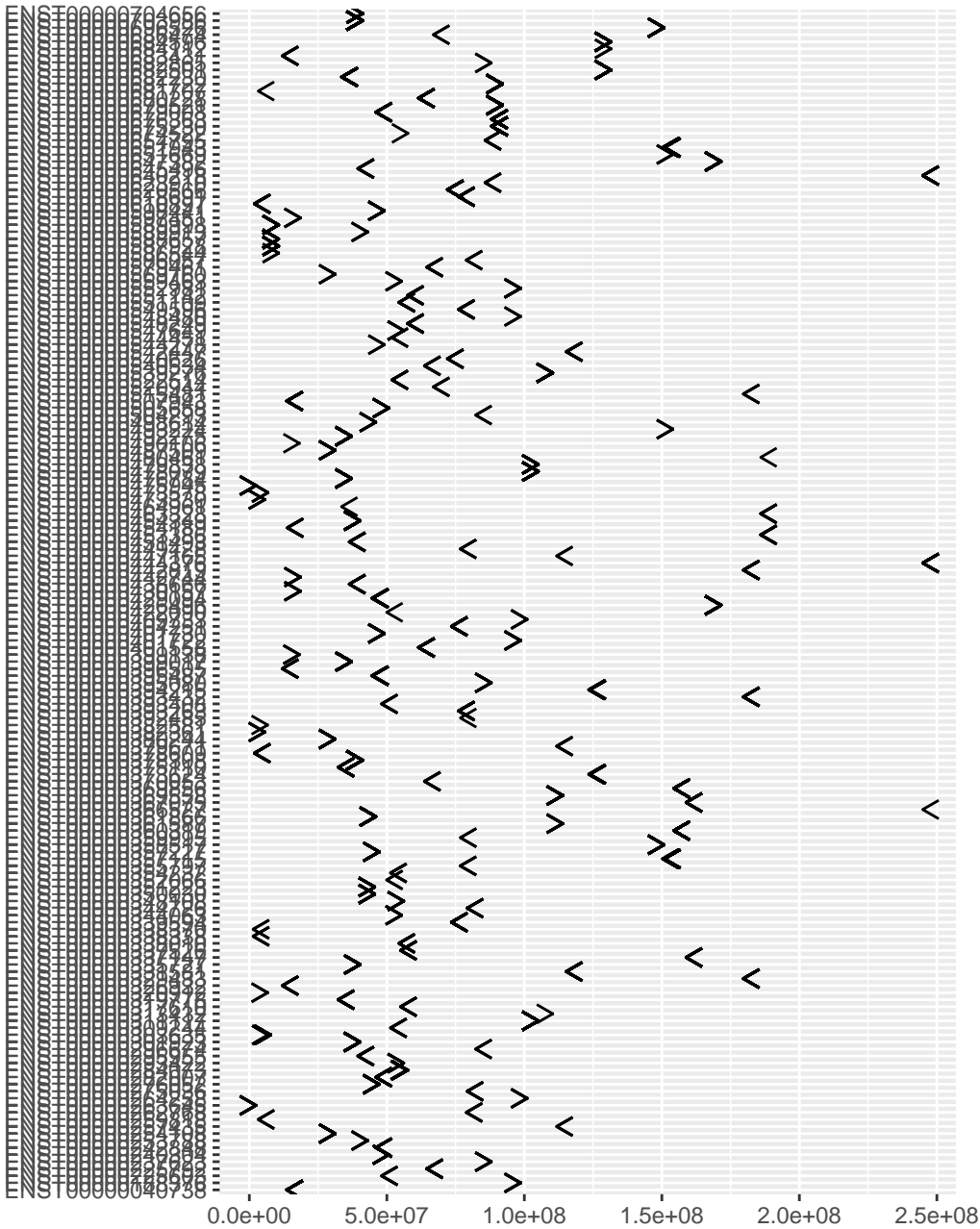
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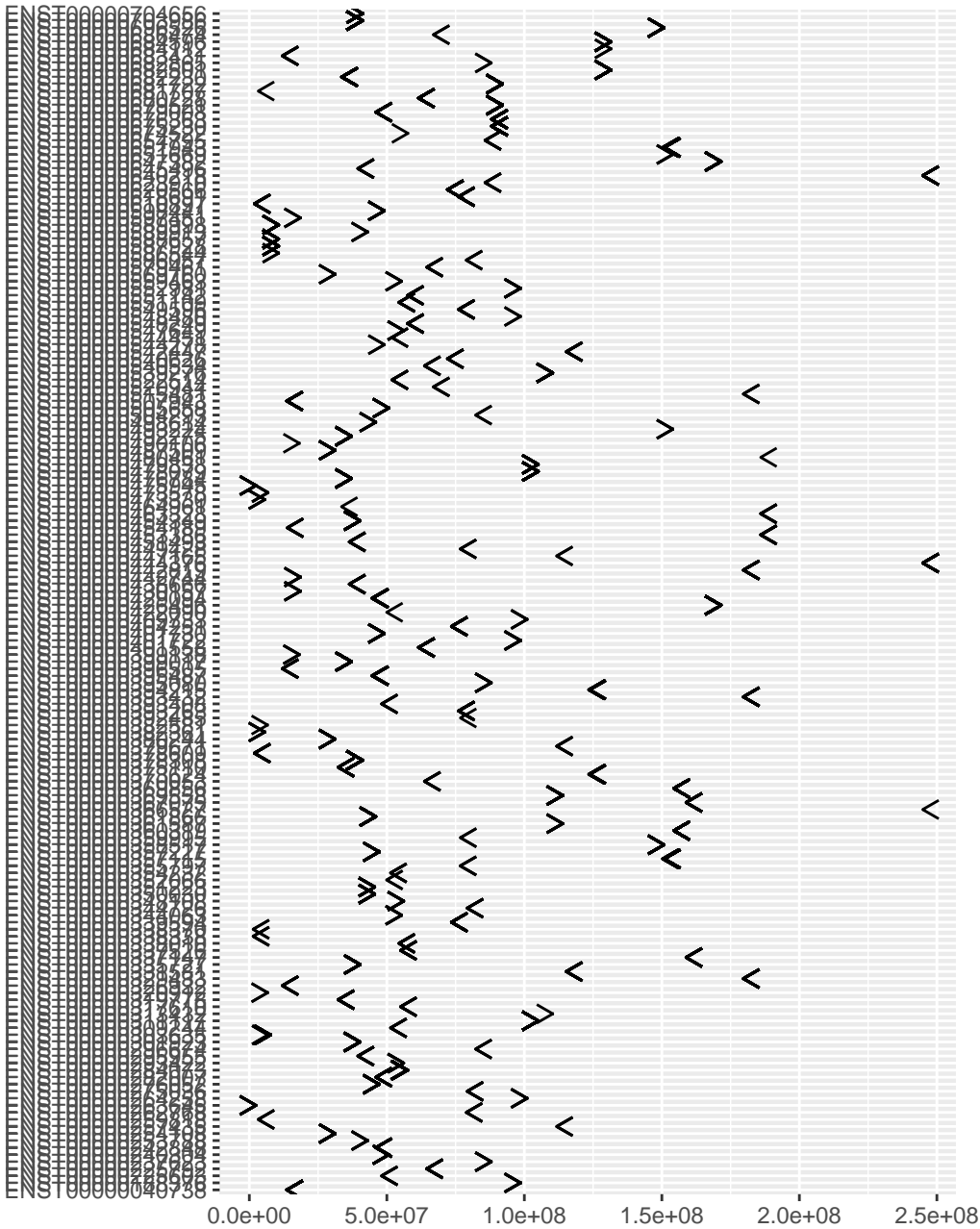


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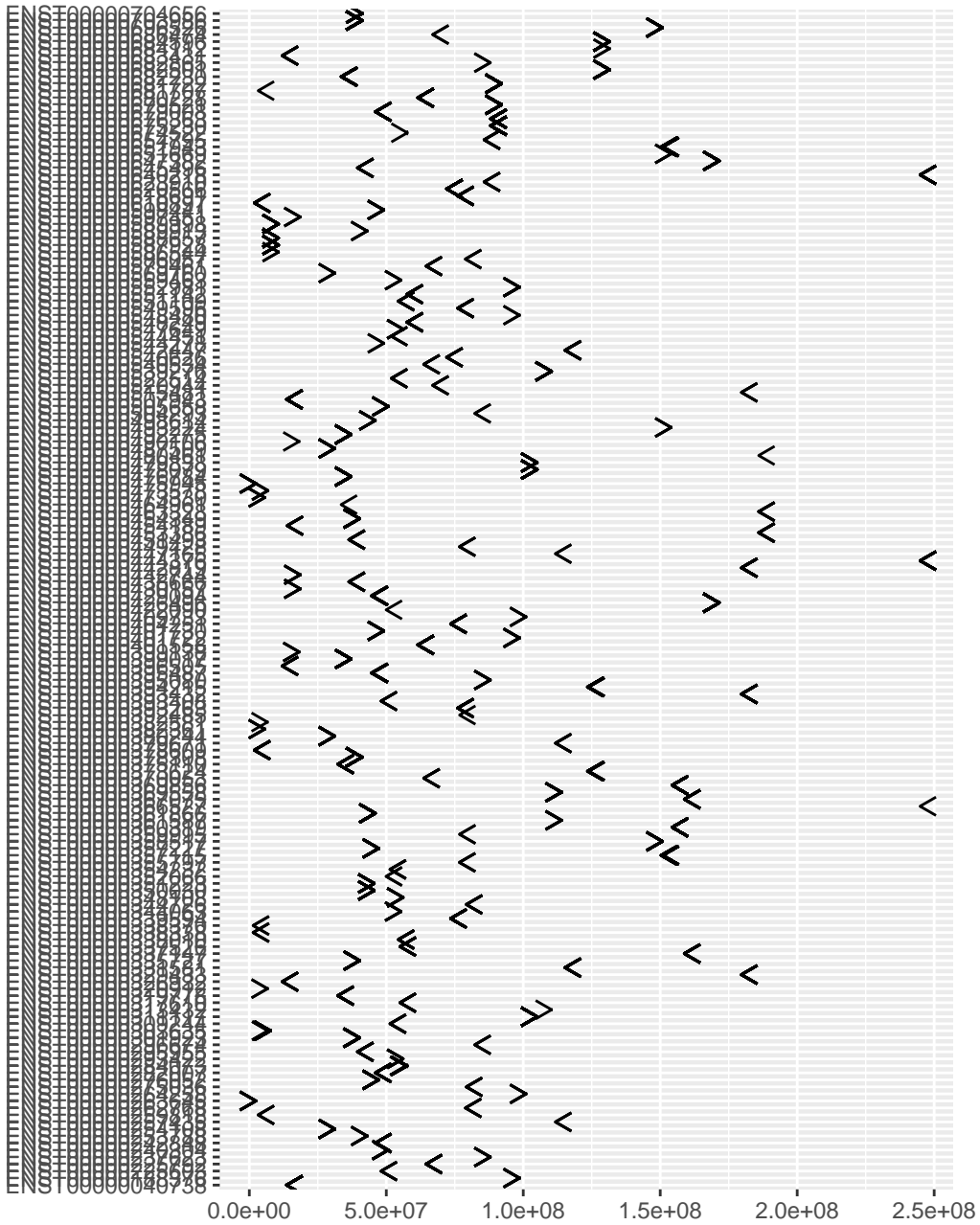


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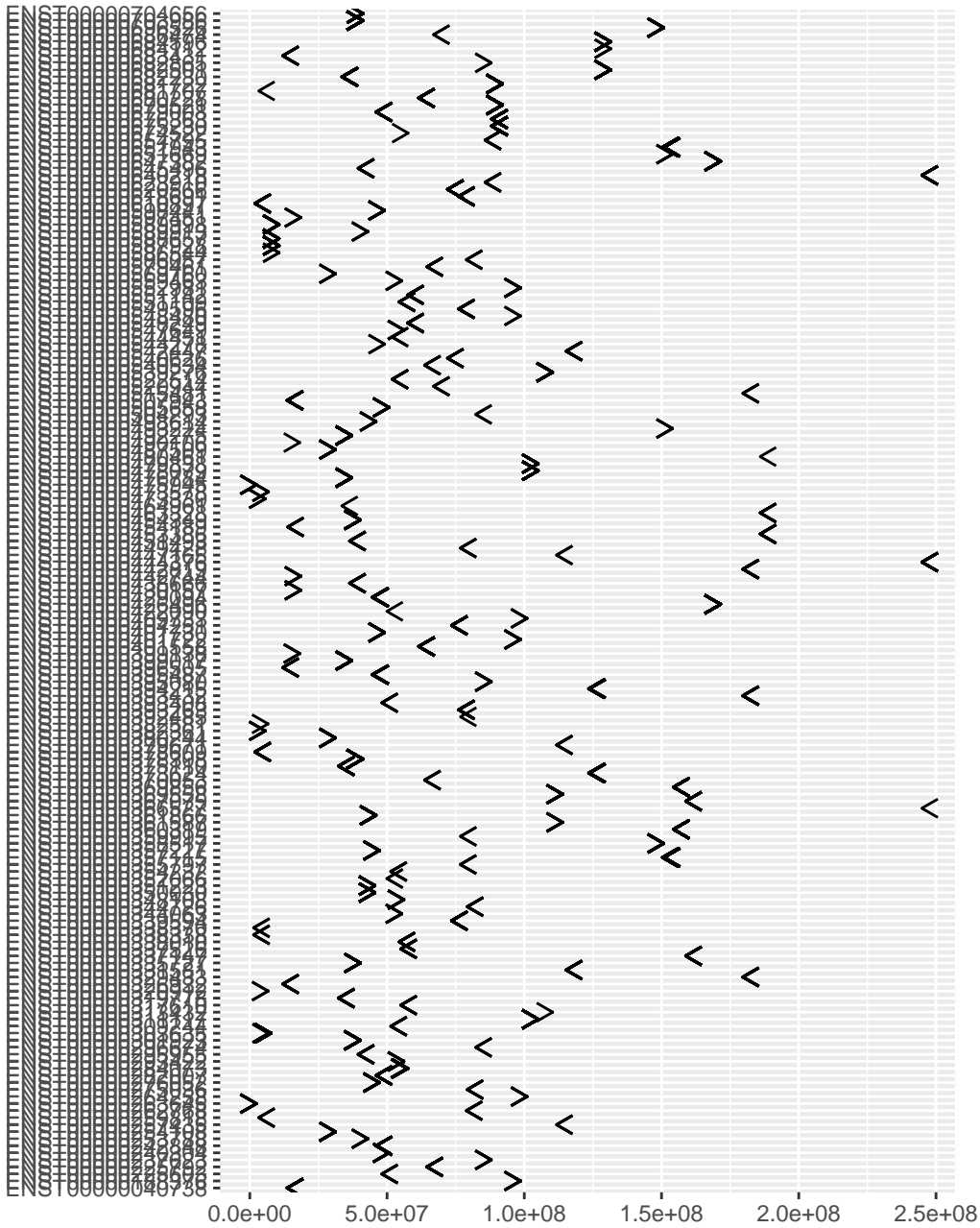
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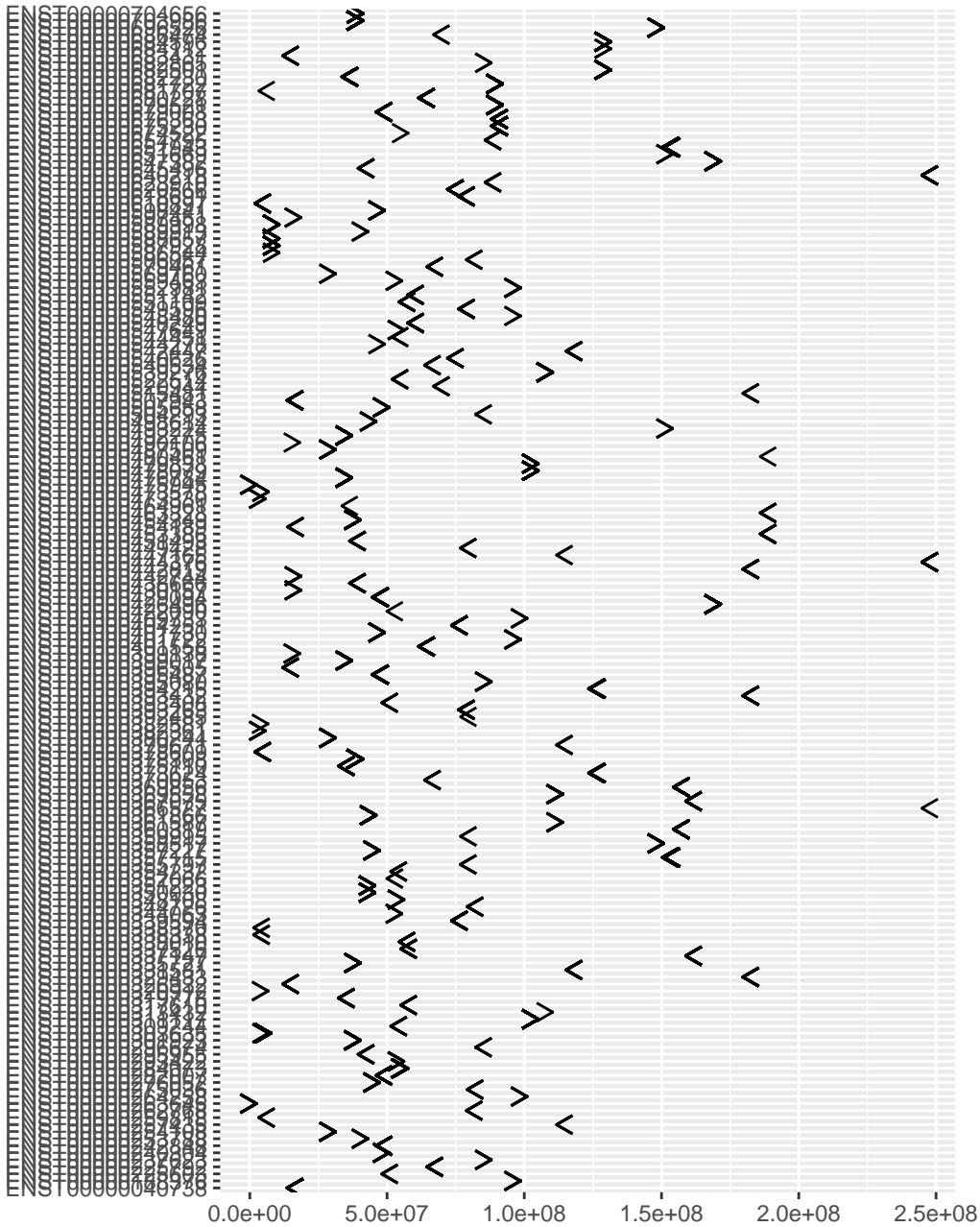
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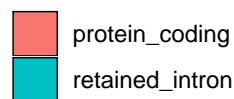
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RPS23

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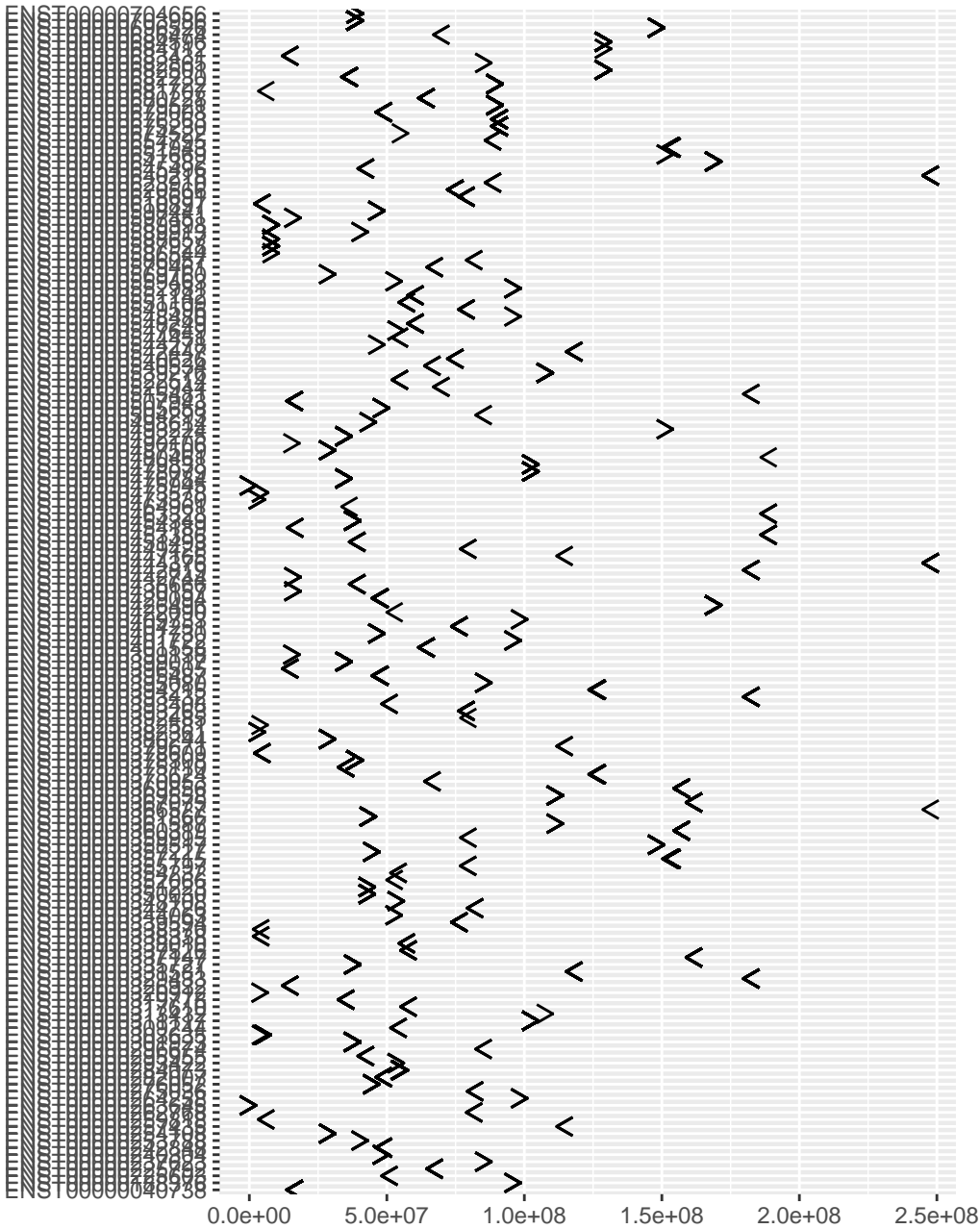


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TNRC6B

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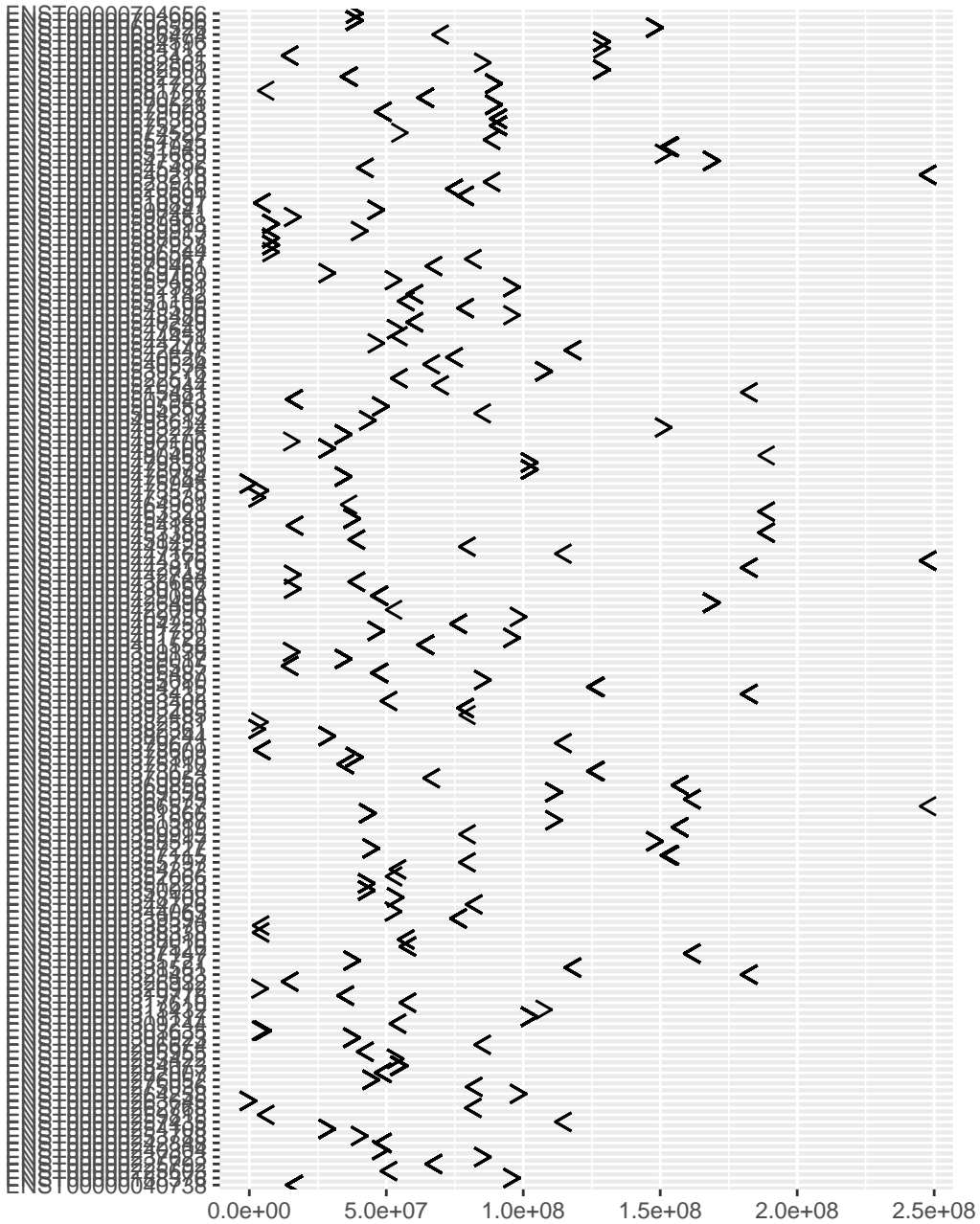
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protein_coding

CAMTA1

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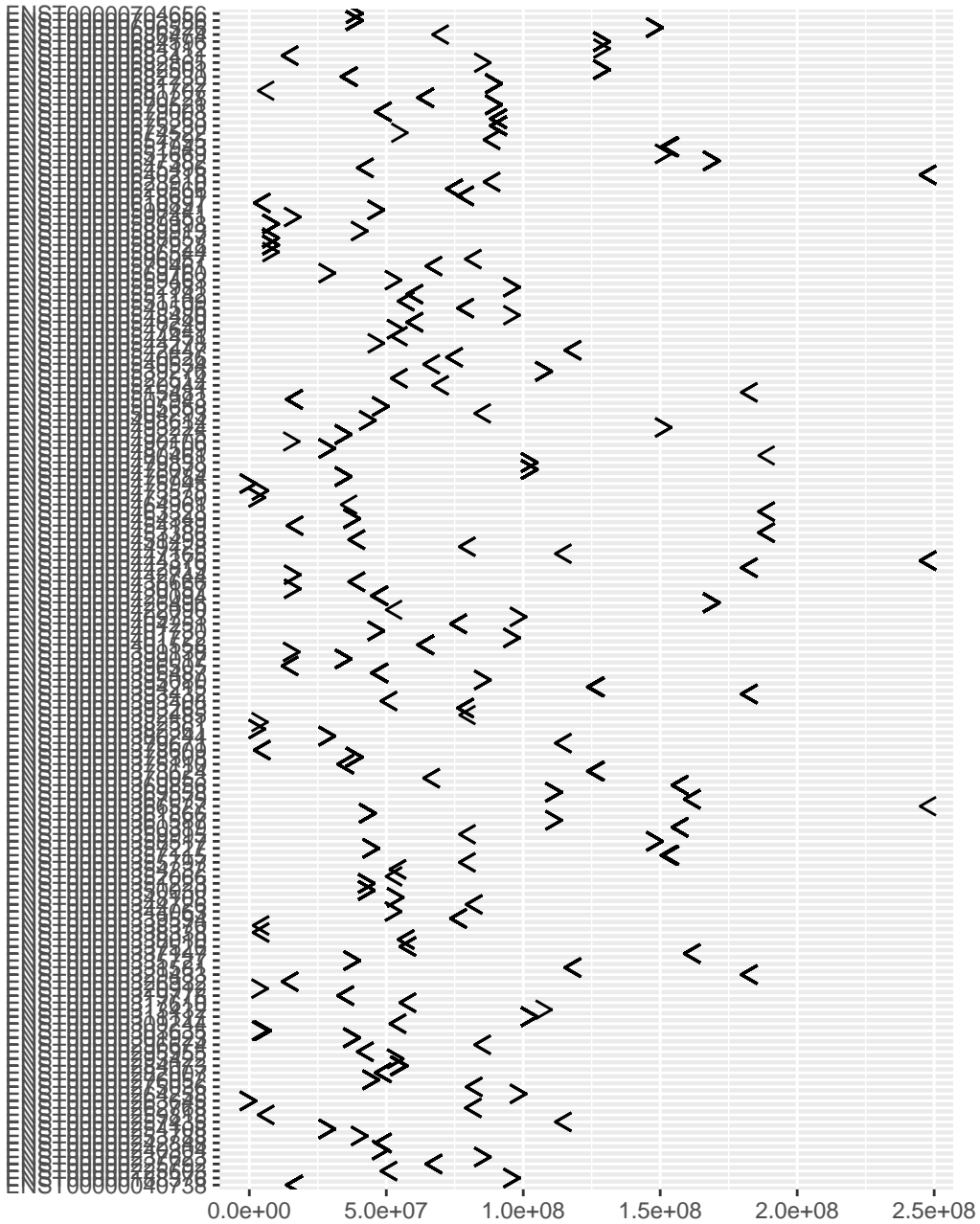
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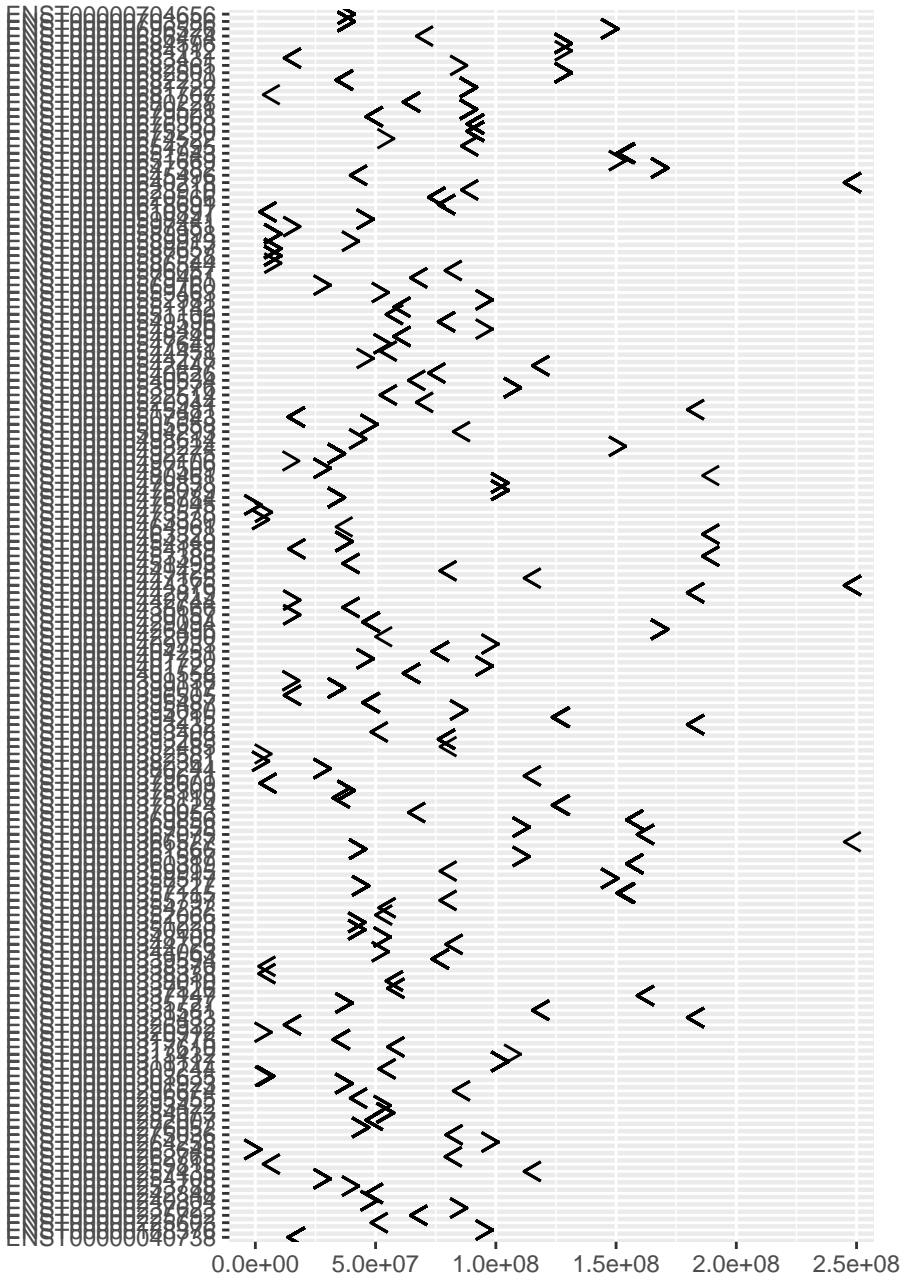
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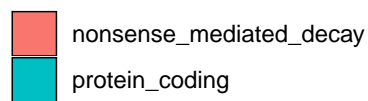


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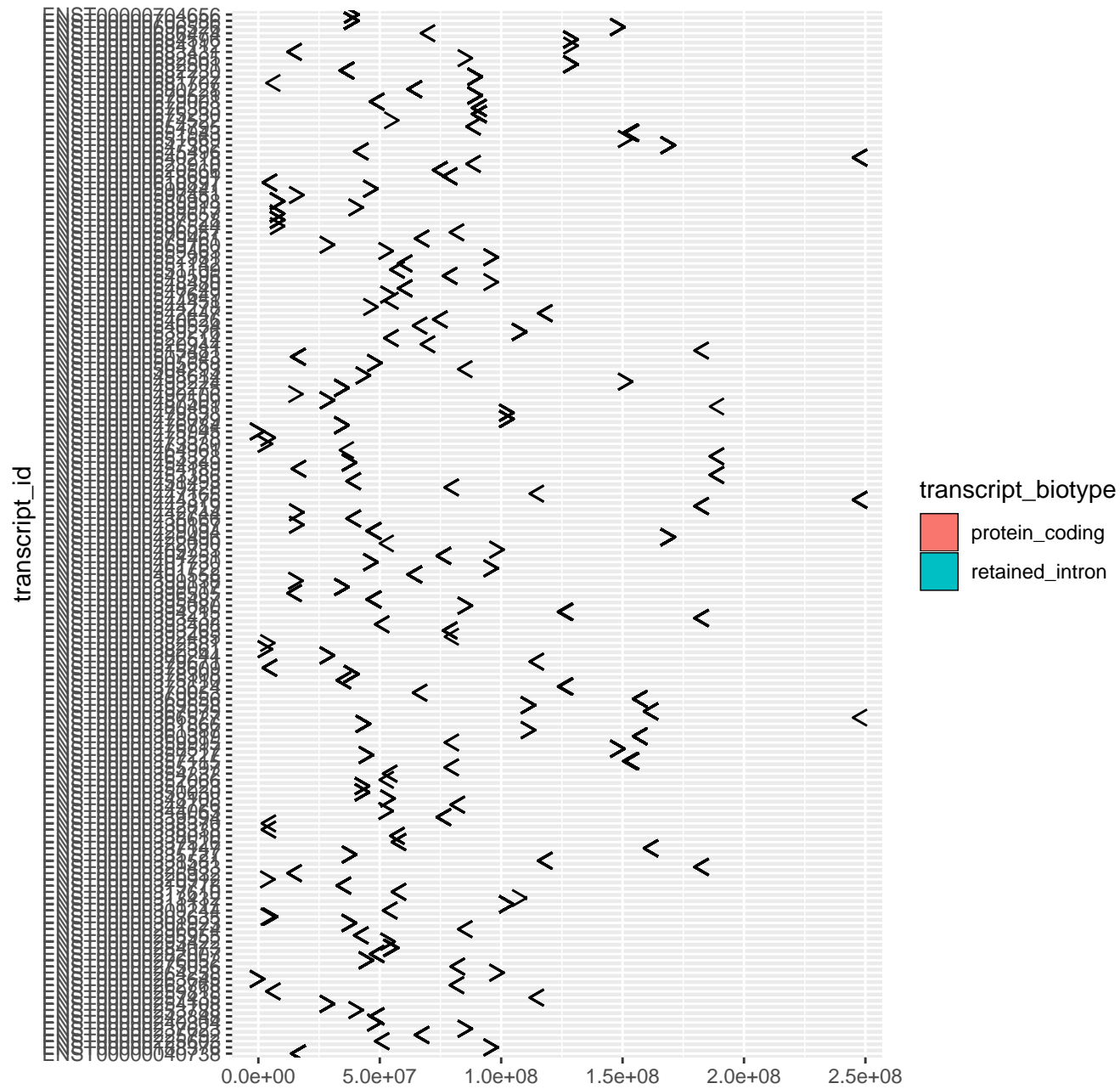
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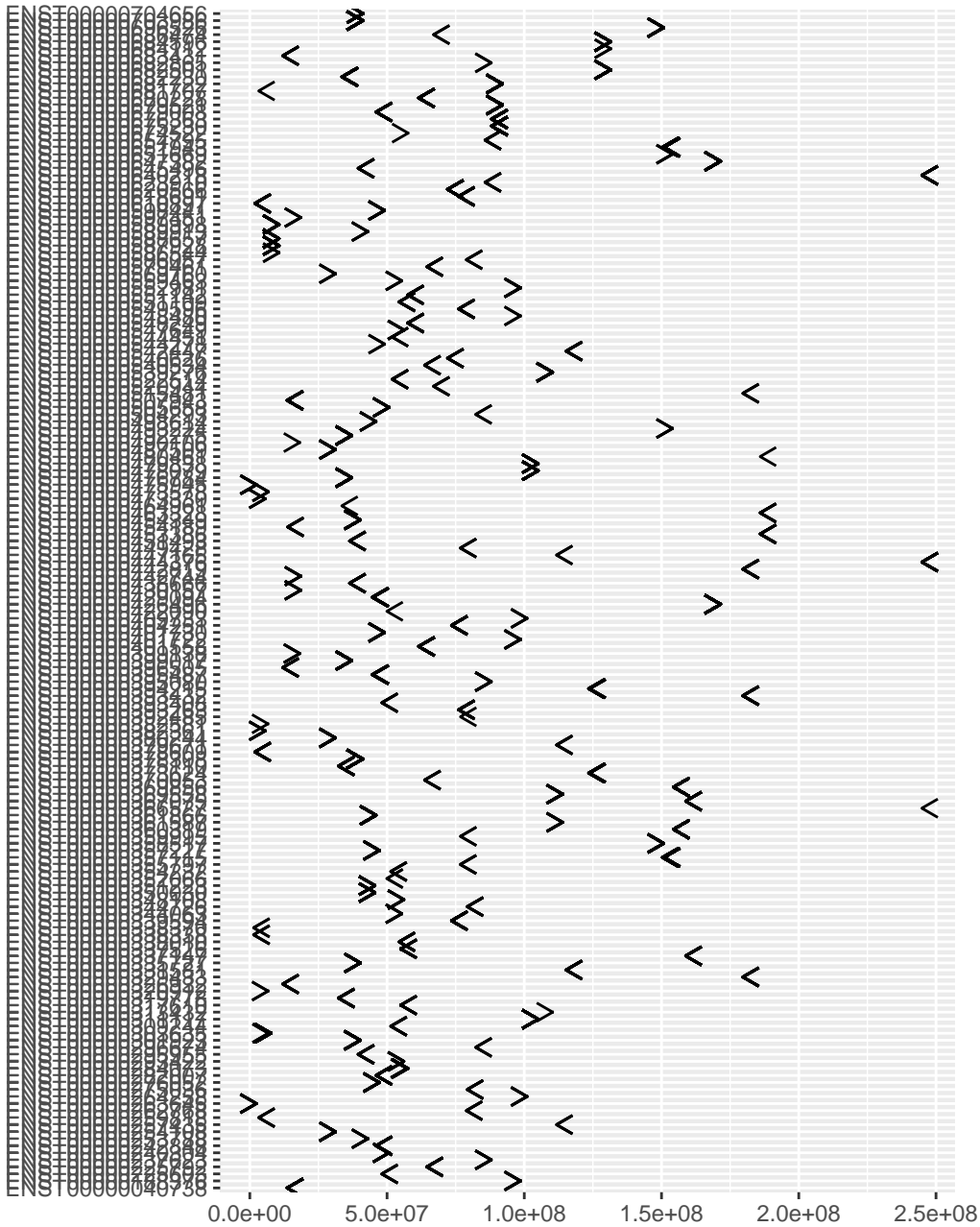


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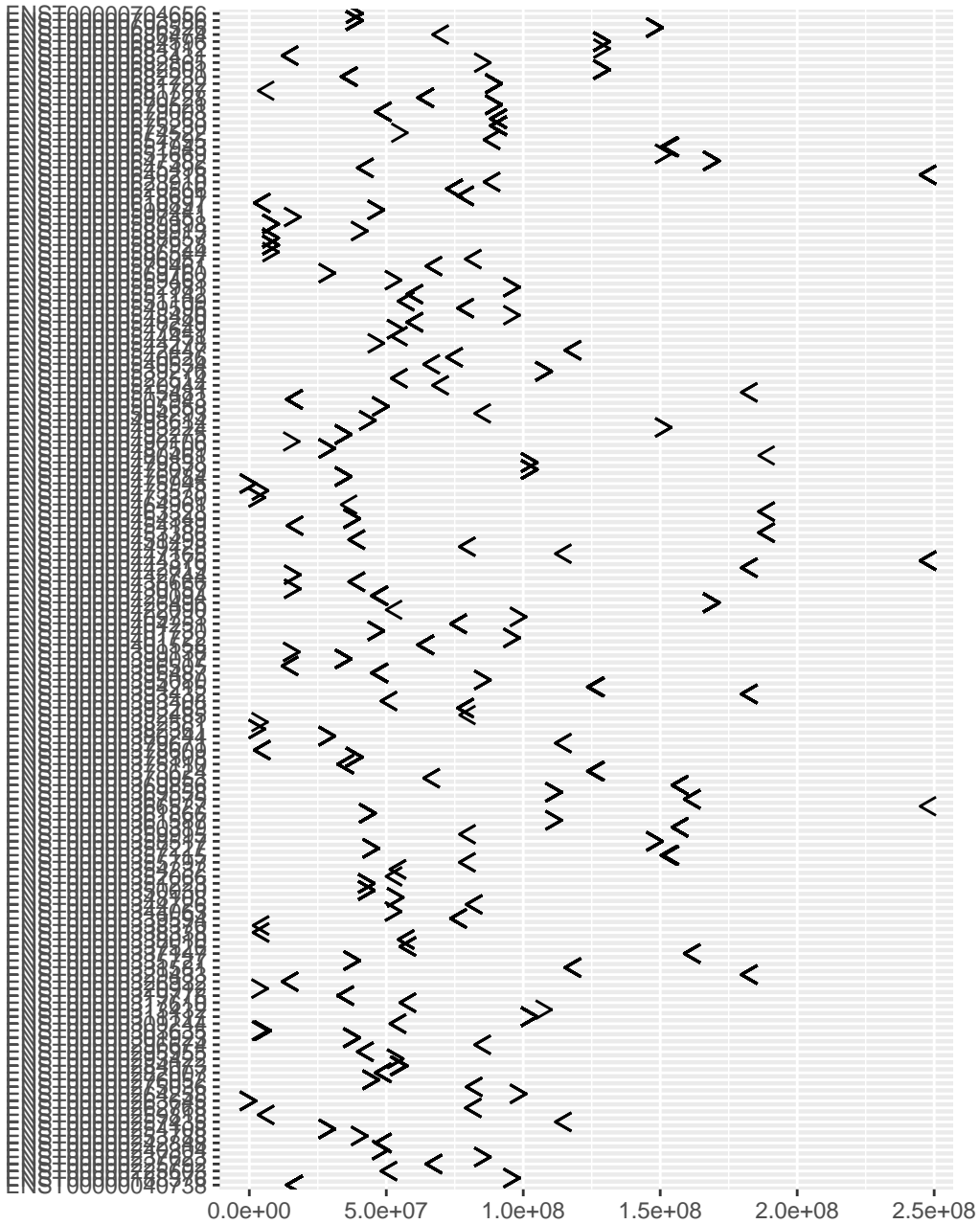
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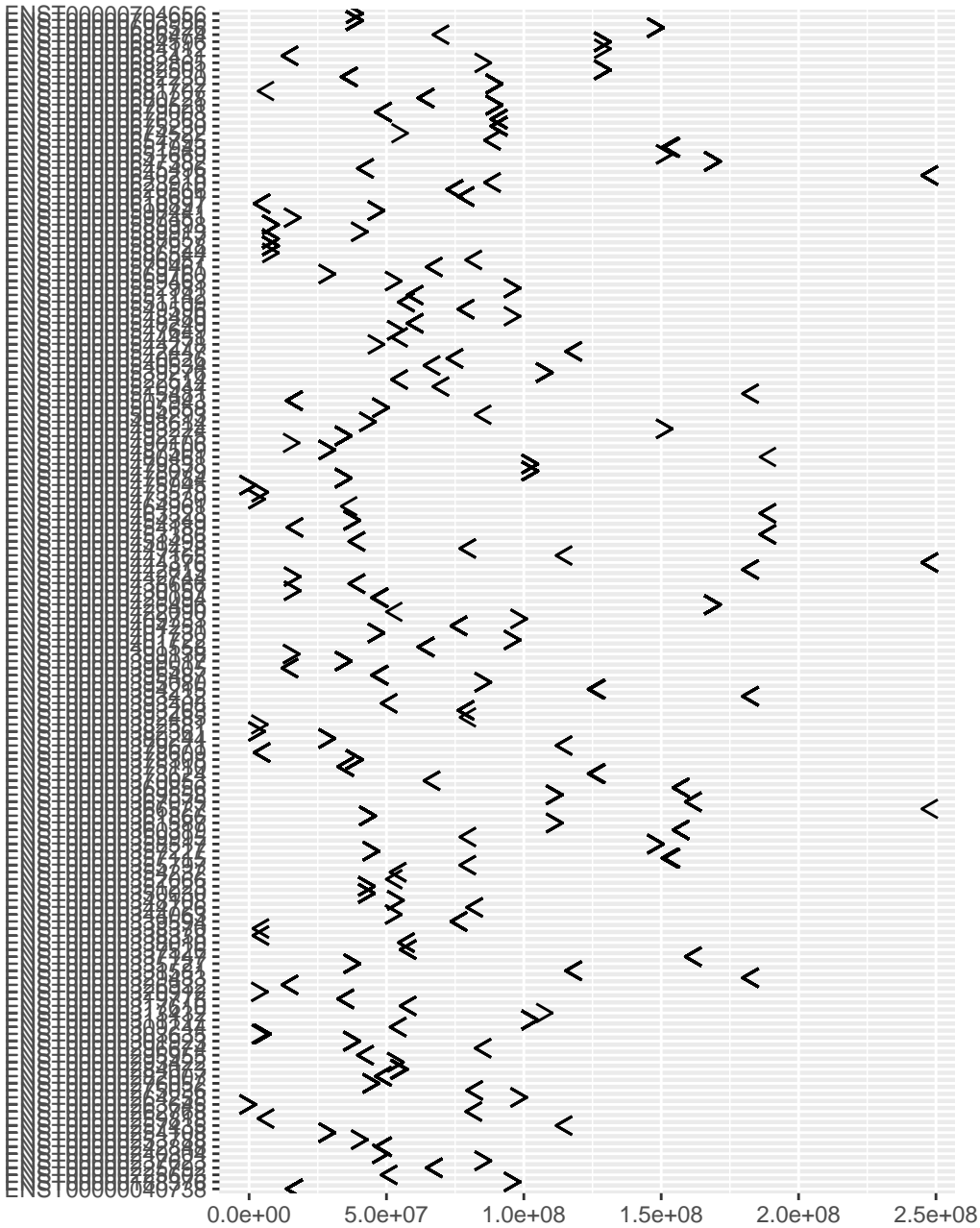
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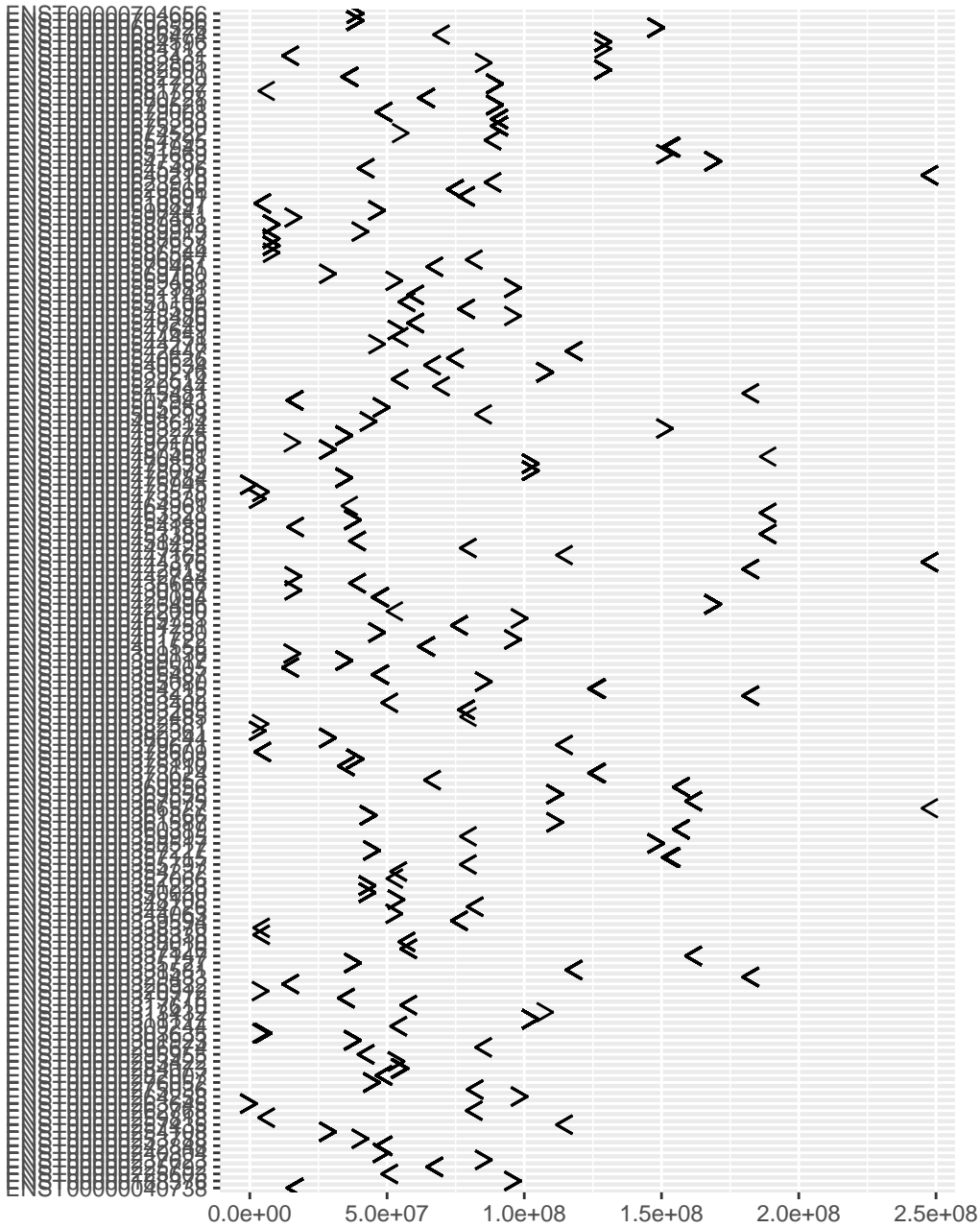
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protein_coding

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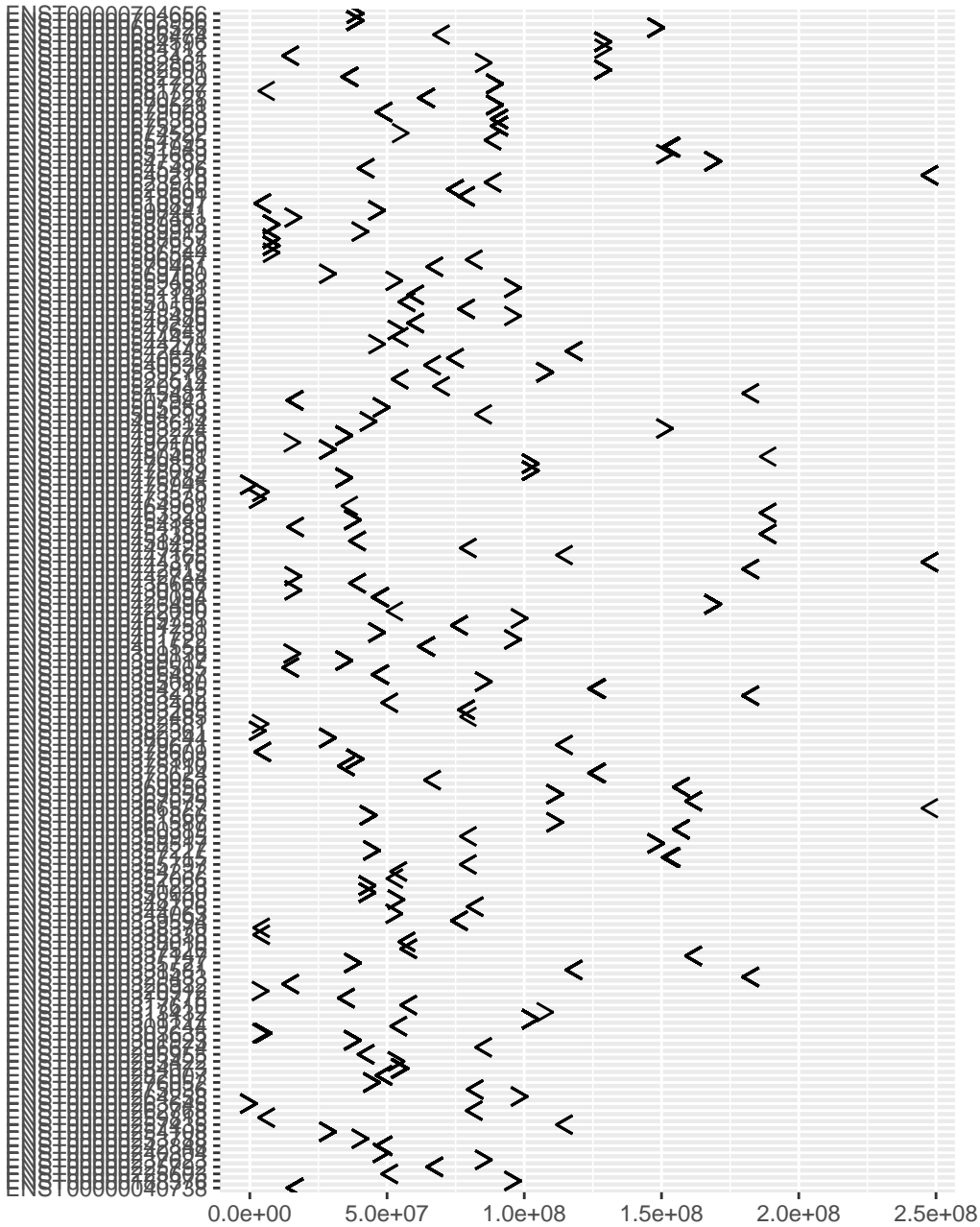


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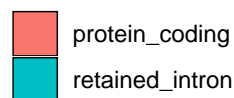
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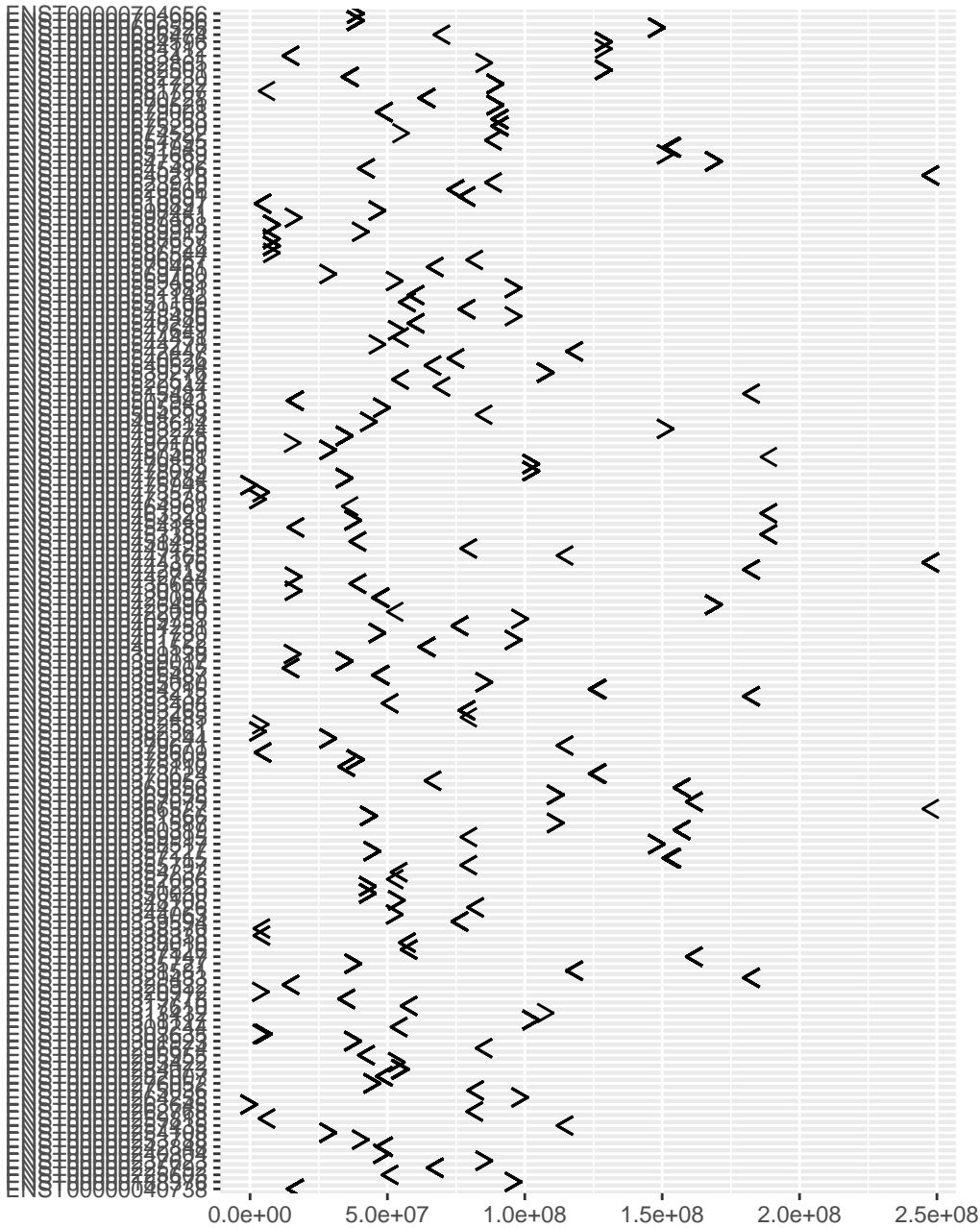


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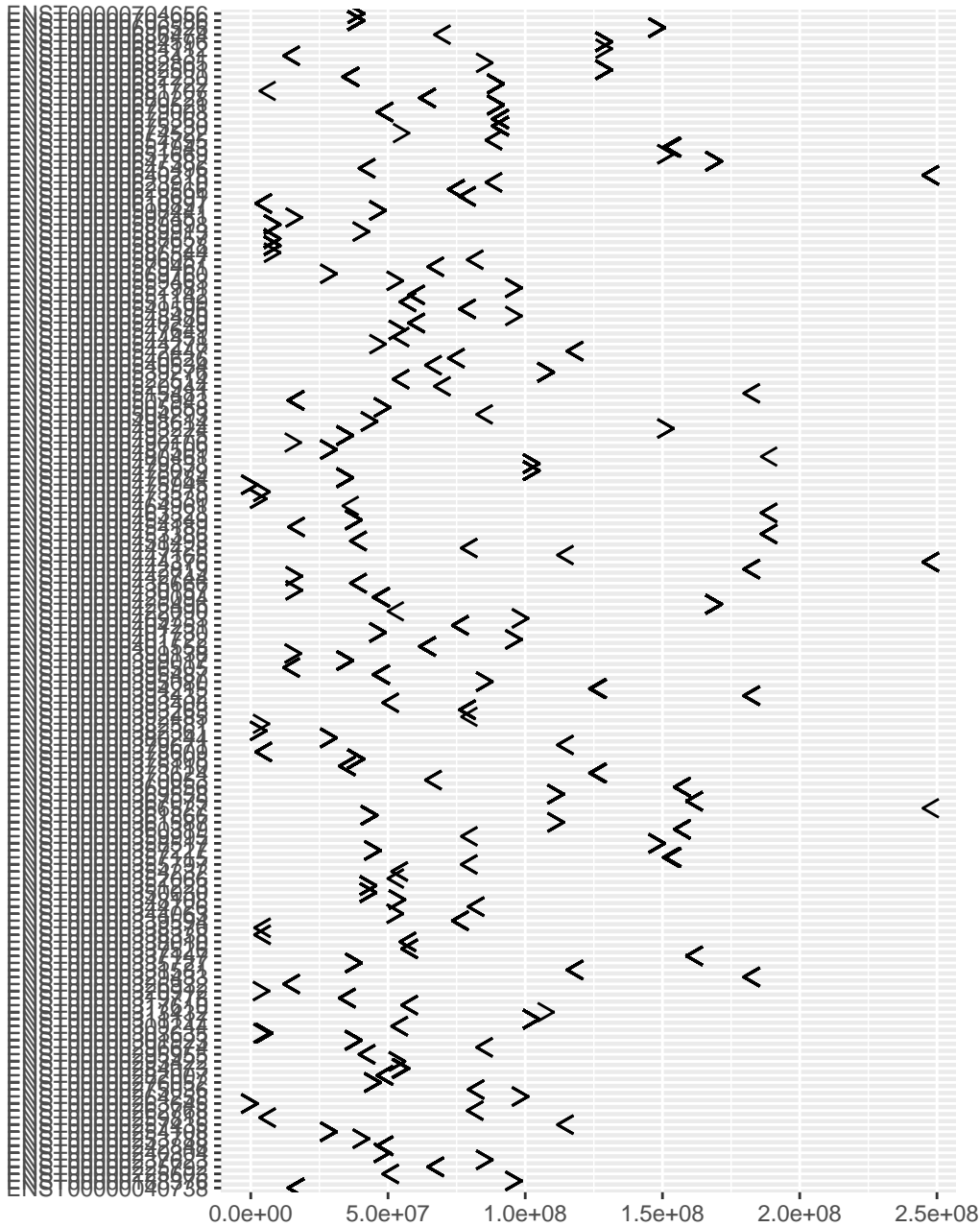
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EZR

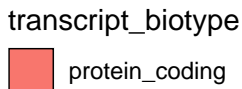
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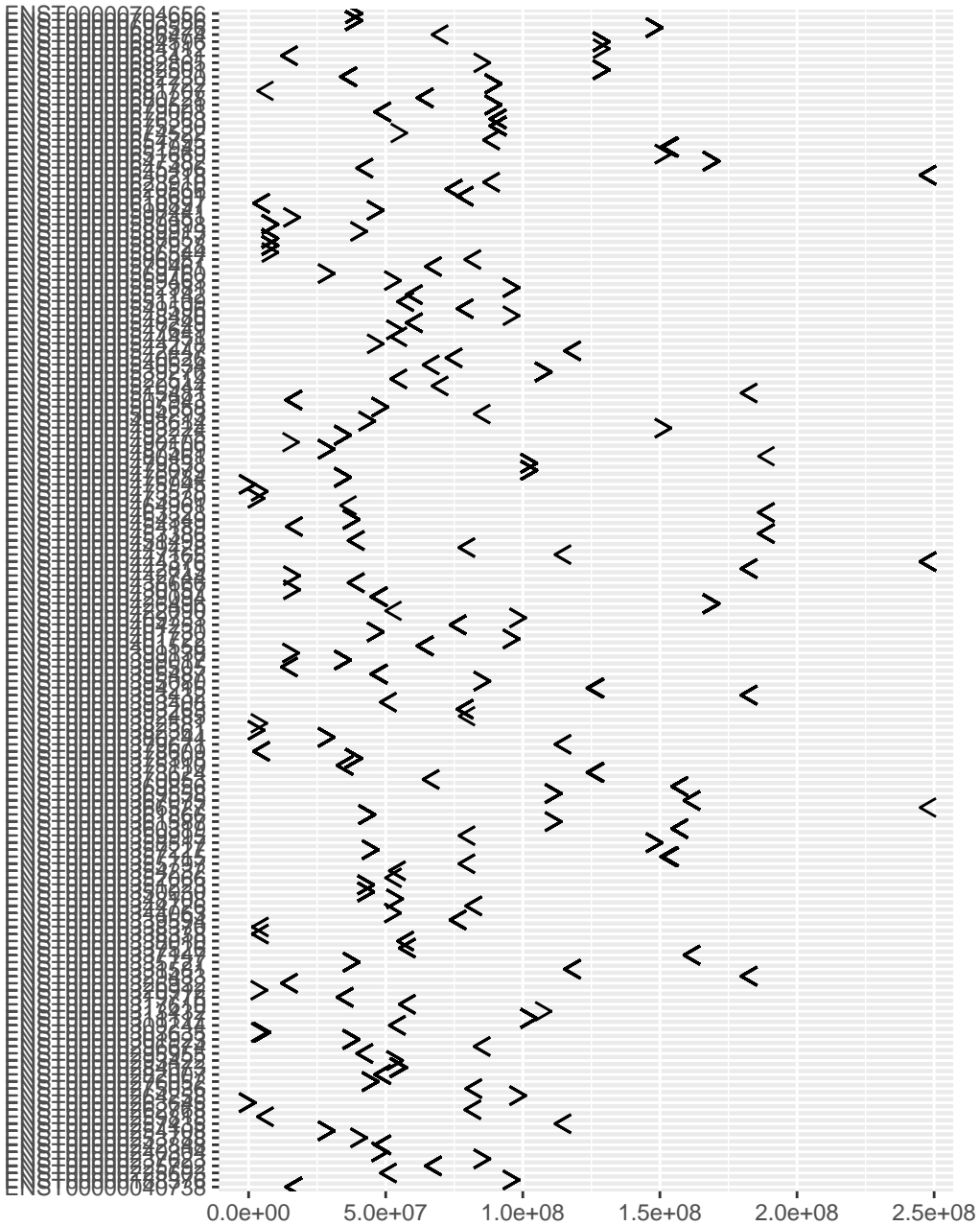
protein_coding

The image displays a highly complex and dense pattern, likely a digital artwork or a data visualization. It features a vast array of small, stylized, and somewhat abstract shapes, possibly representing a complex data set or a highly detailed texture. The pattern is composed of many small, interconnected elements that form a larger, intricate structure. The overall appearance is that of a highly detailed, almost fractal-like design, with a strong sense of repetition and symmetry. The pattern is rendered in a monochromatic style, using shades of gray and black on a white background. The elements are arranged in a way that creates a sense of depth and movement, with some areas appearing more prominent than others. The overall effect is one of a highly detailed and complex visual structure, possibly representing a large-scale data set or a highly detailed texture.



AURKAIP1

transcript_id

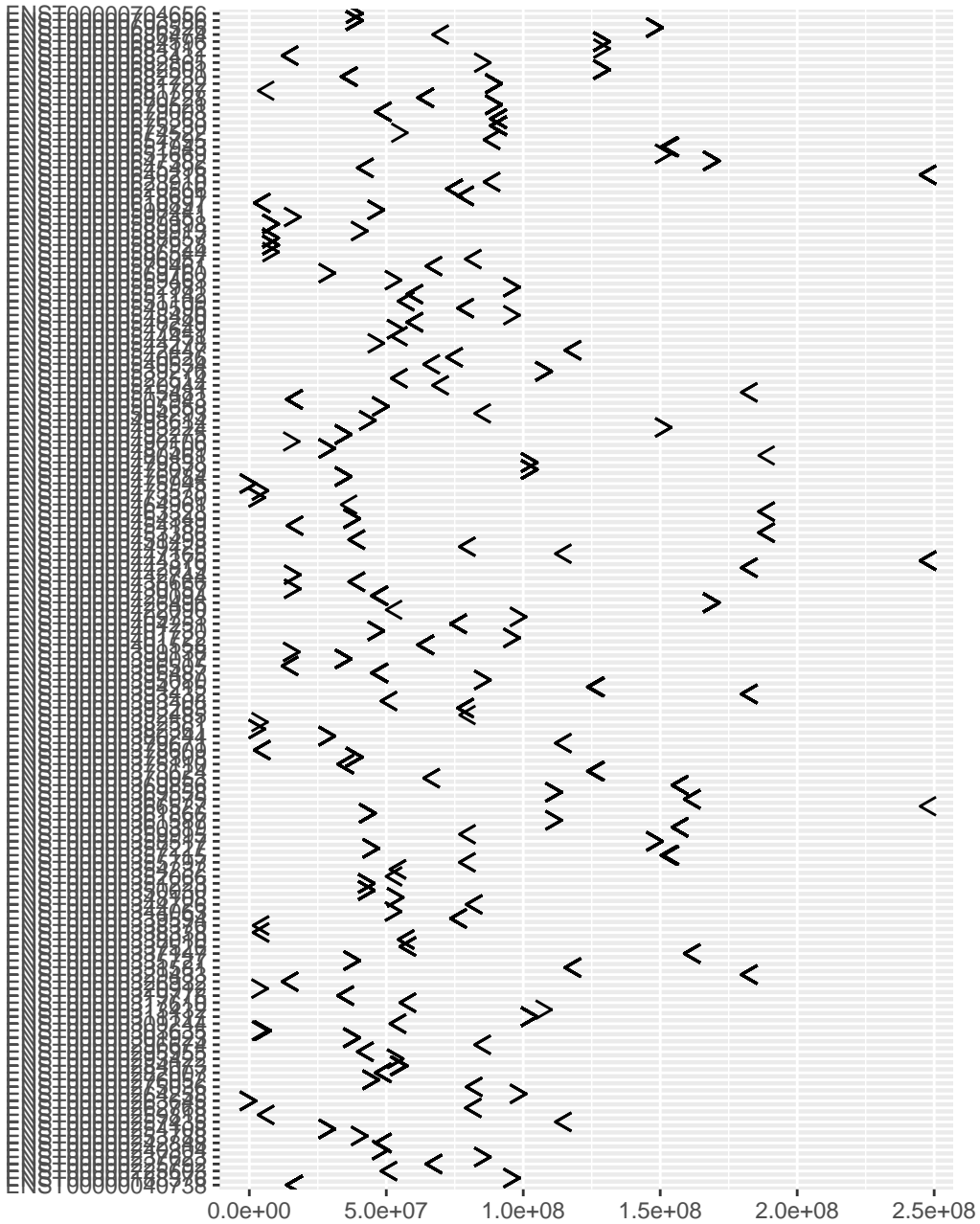


transcript_biotype

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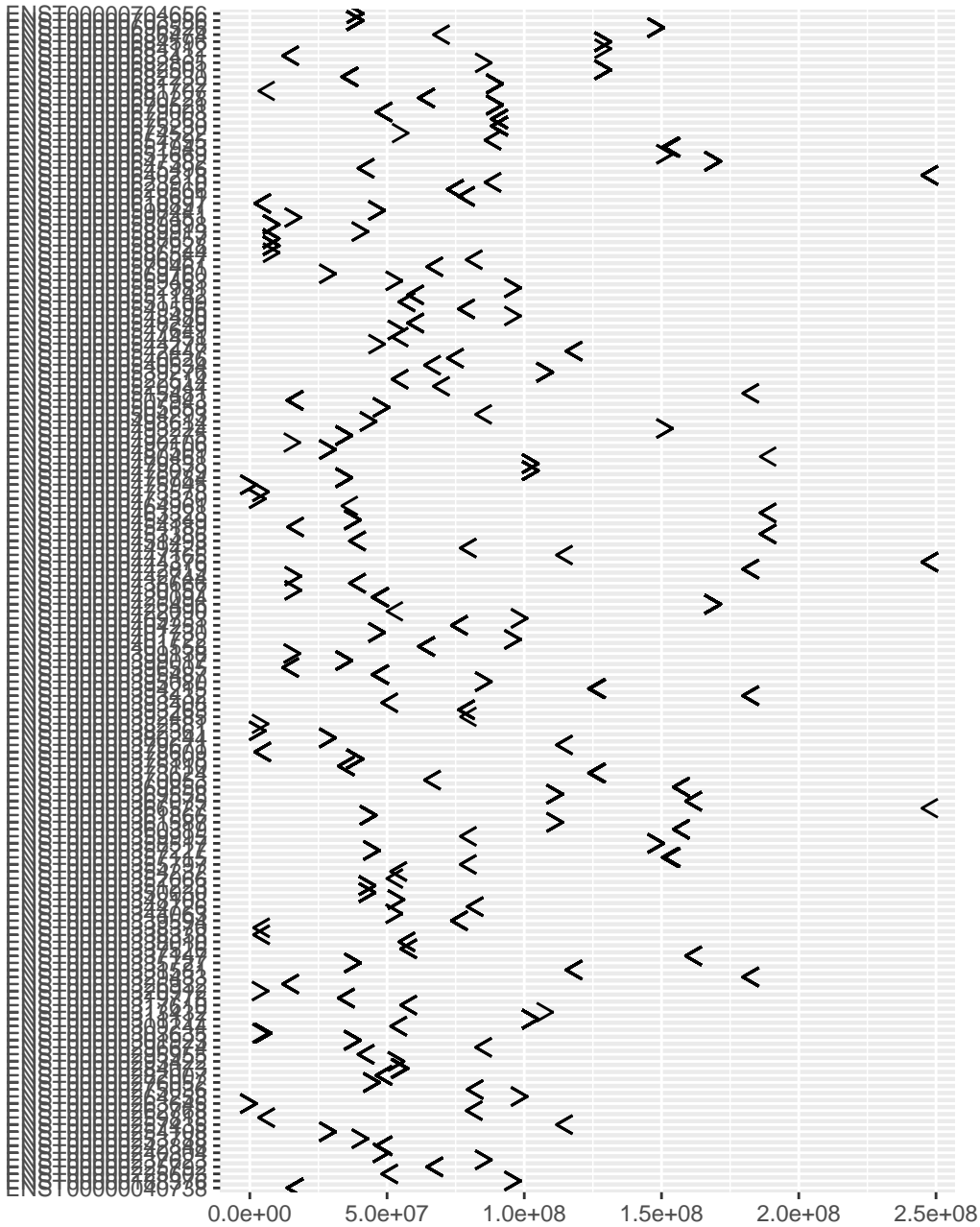
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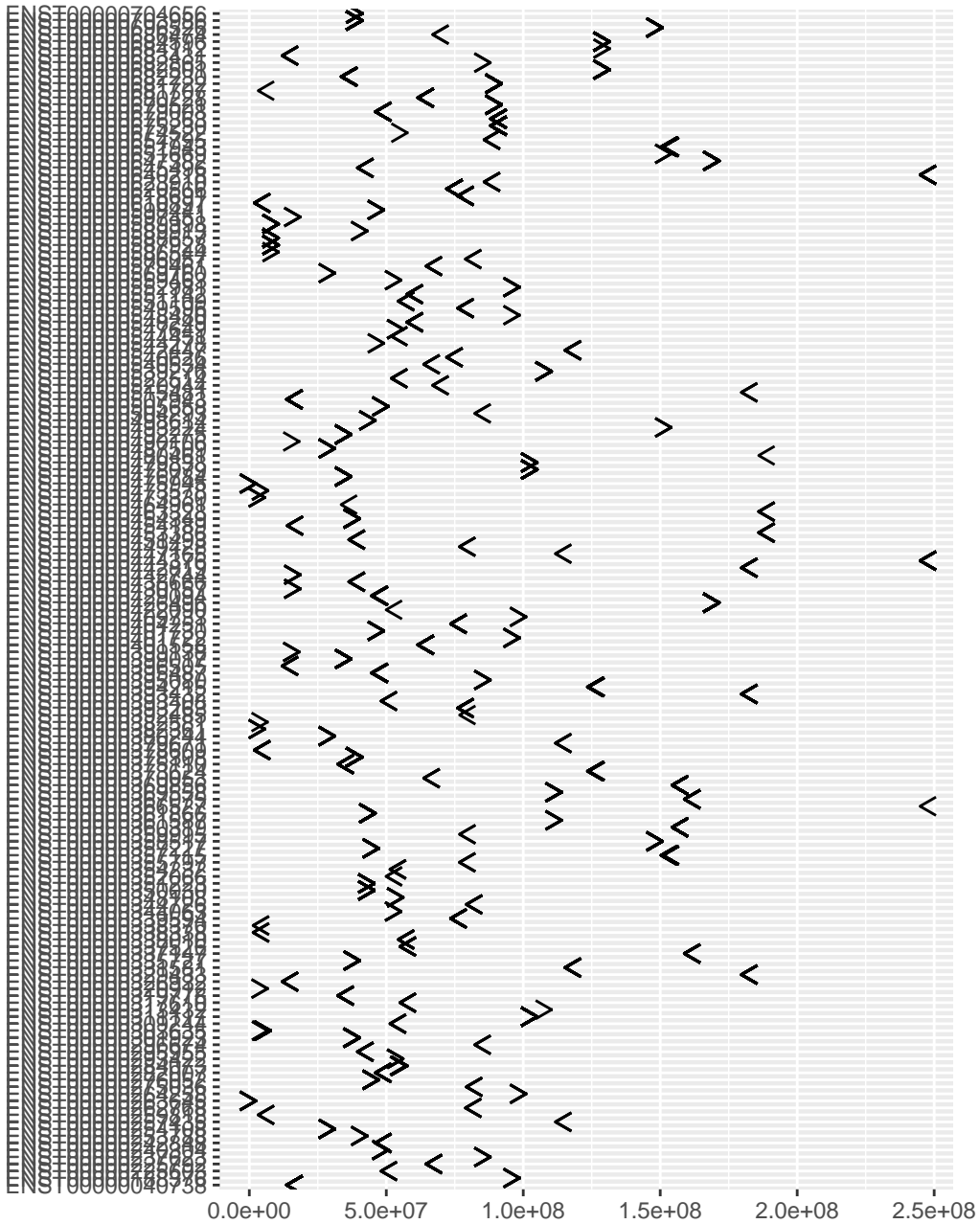
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protein_coding

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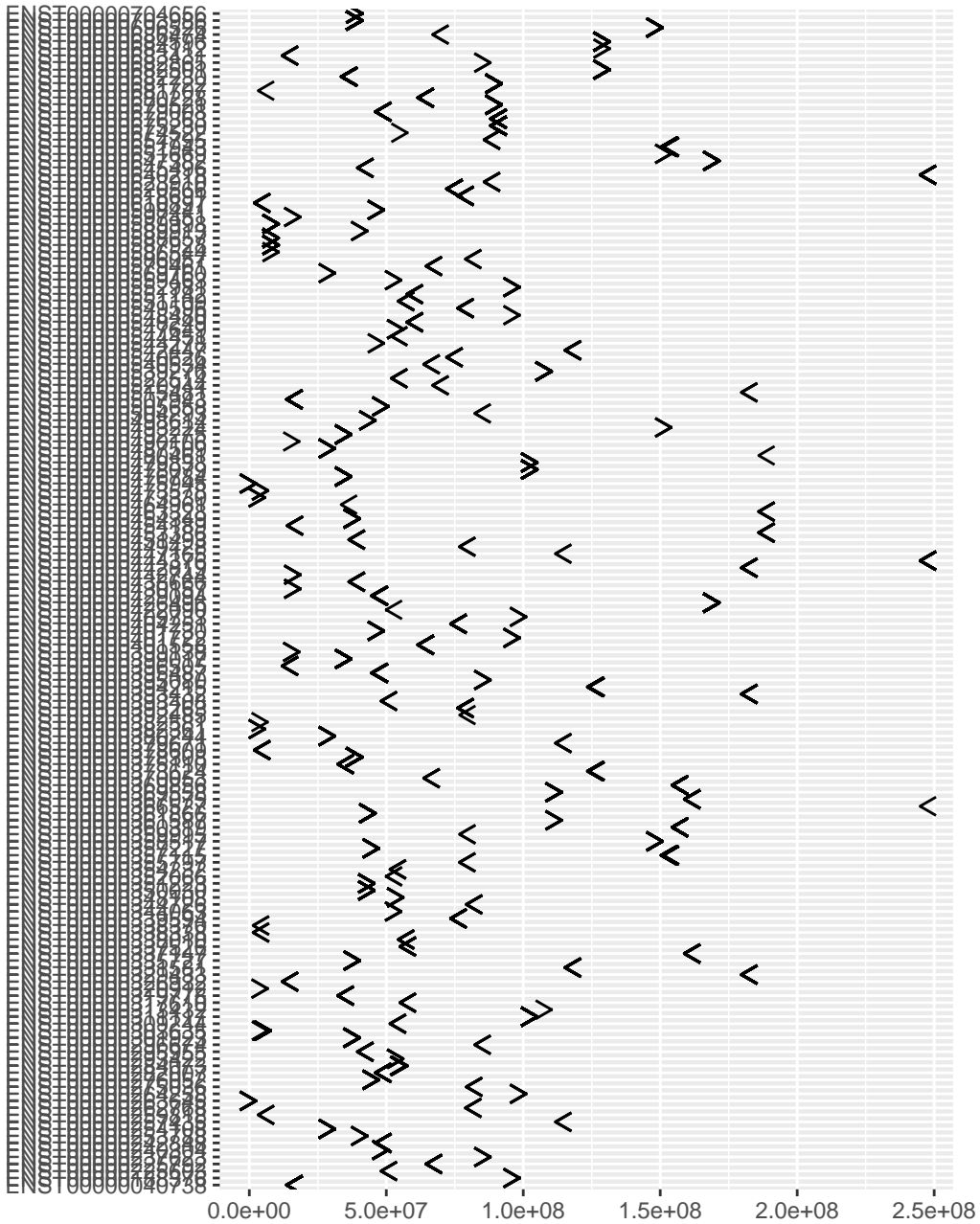
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protein_coding

NUP62

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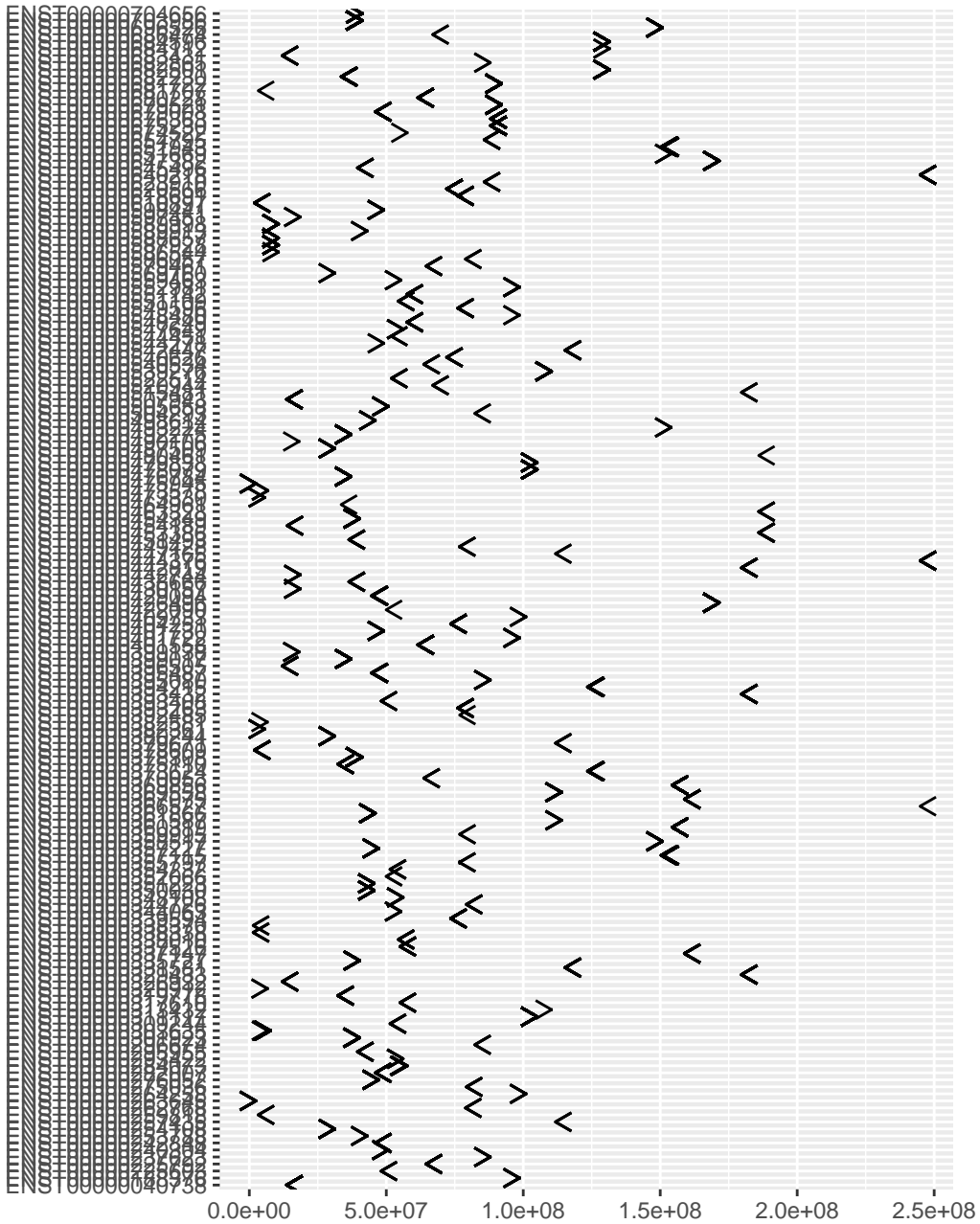
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protein_coding

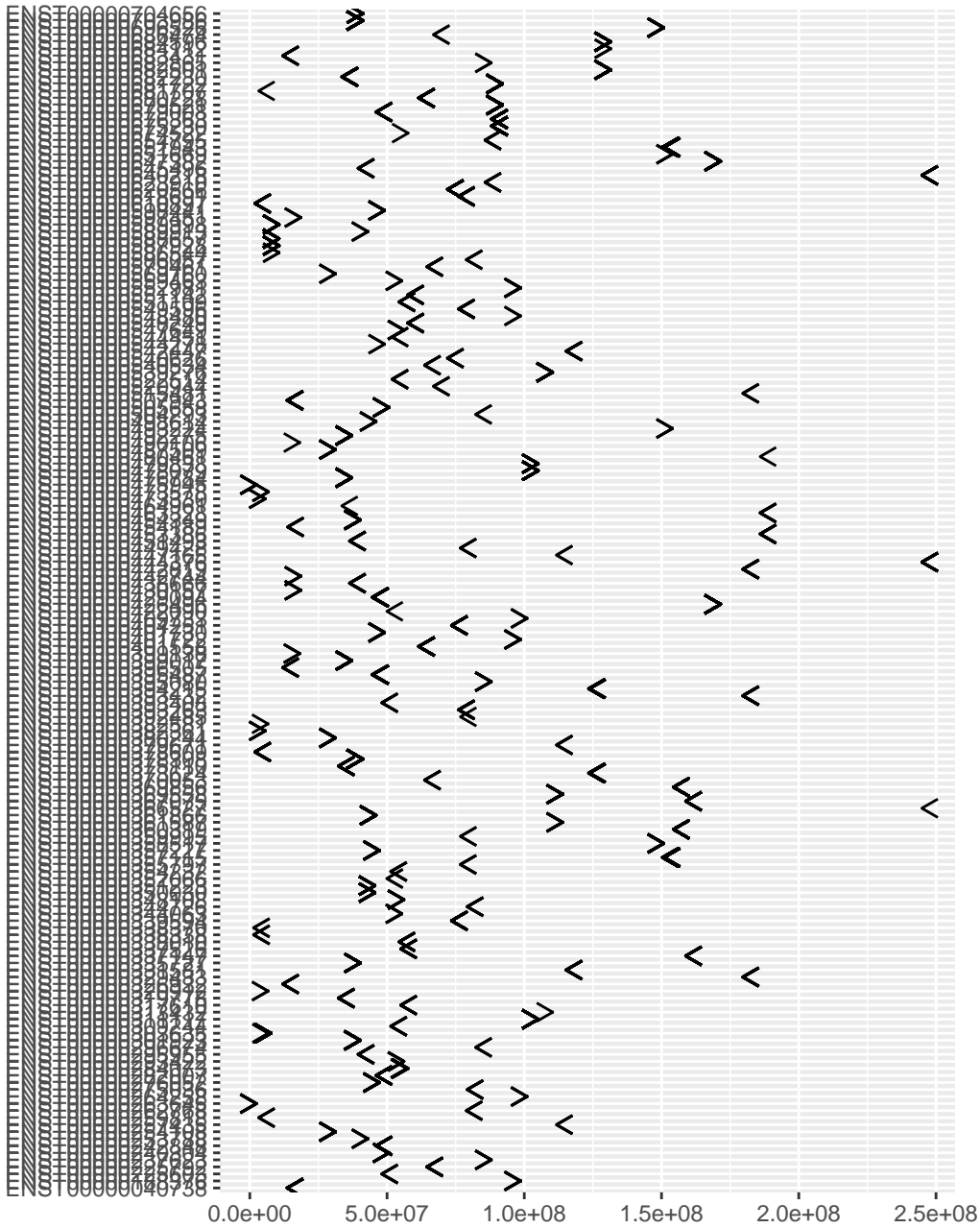
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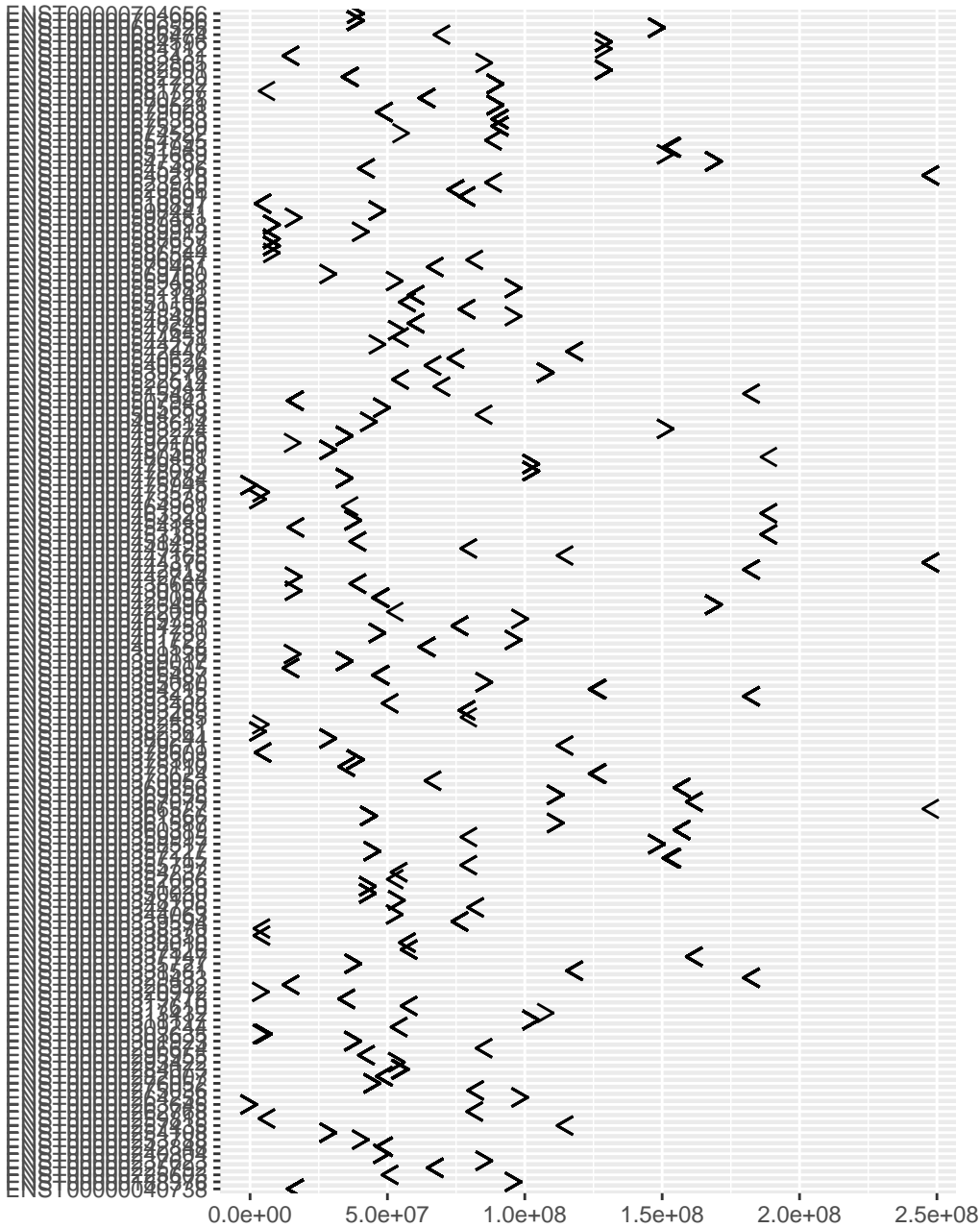
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protein_coding

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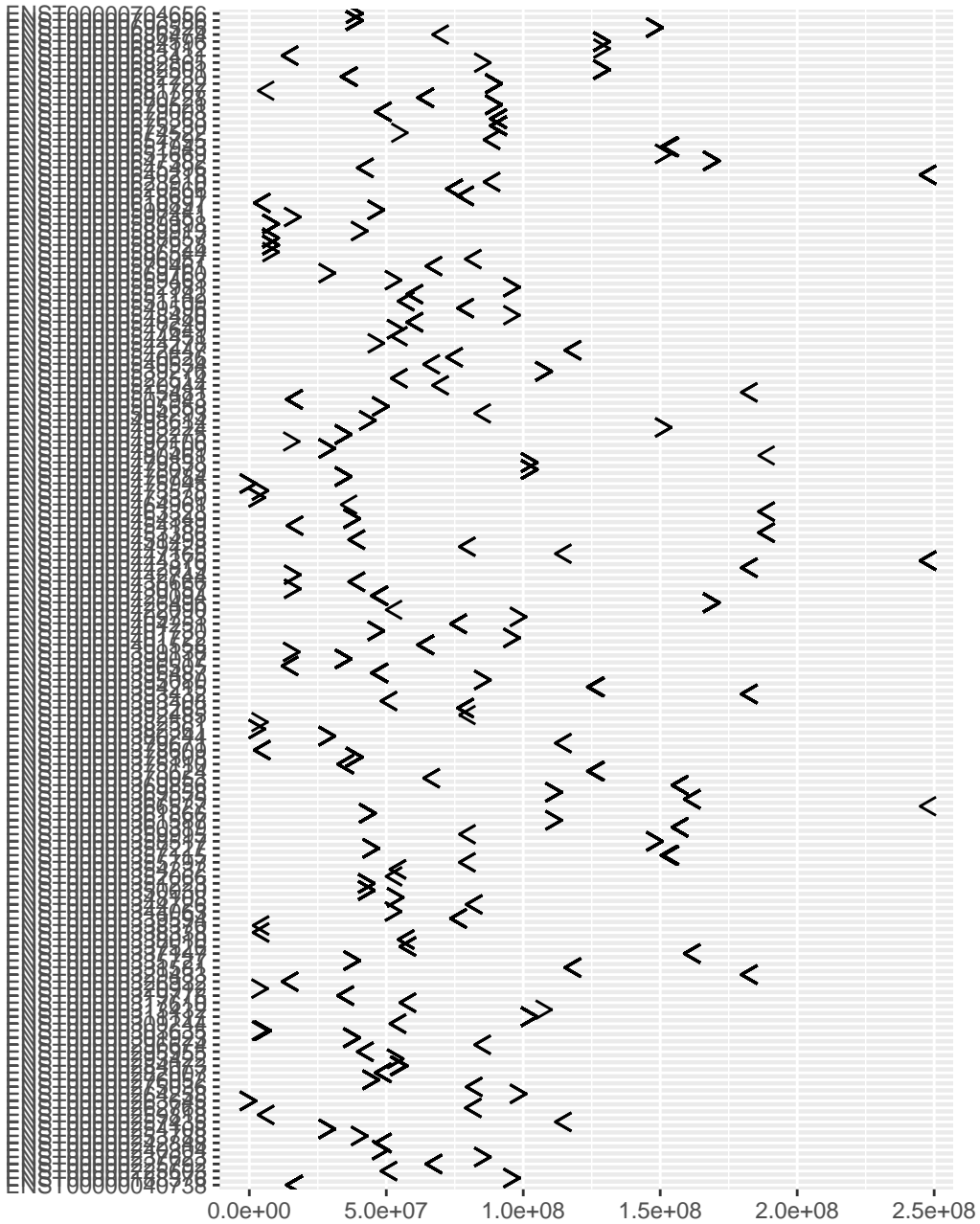
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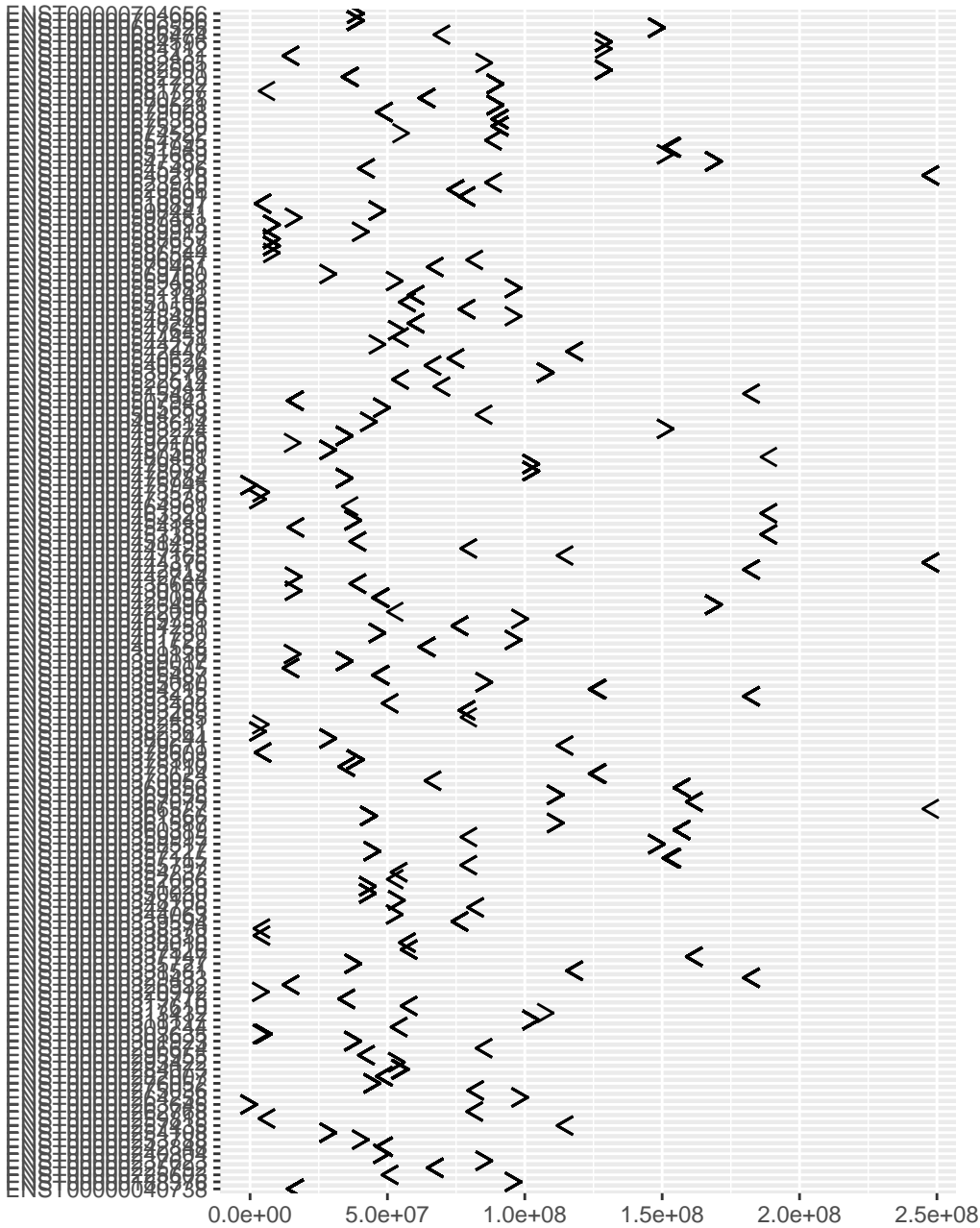
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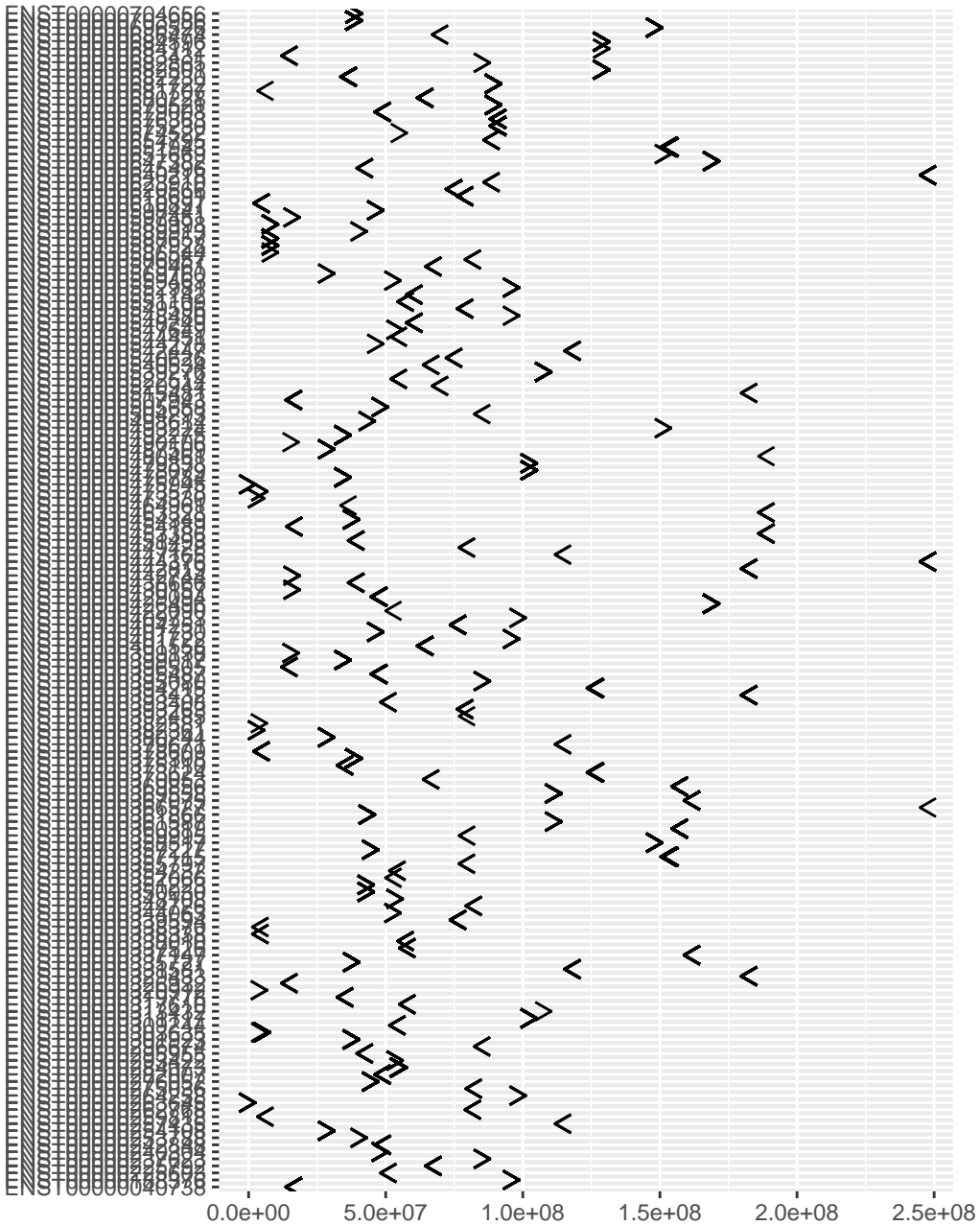
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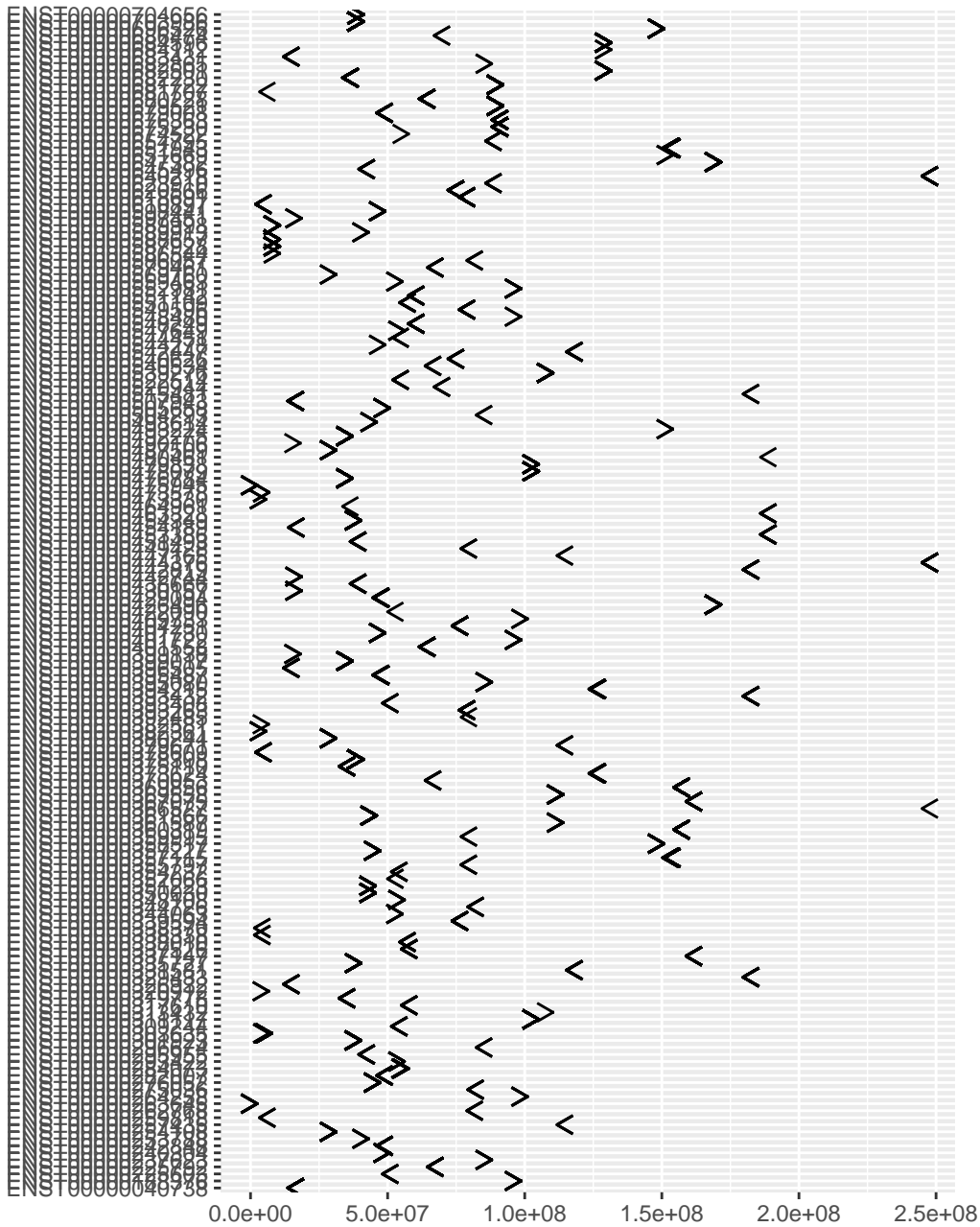


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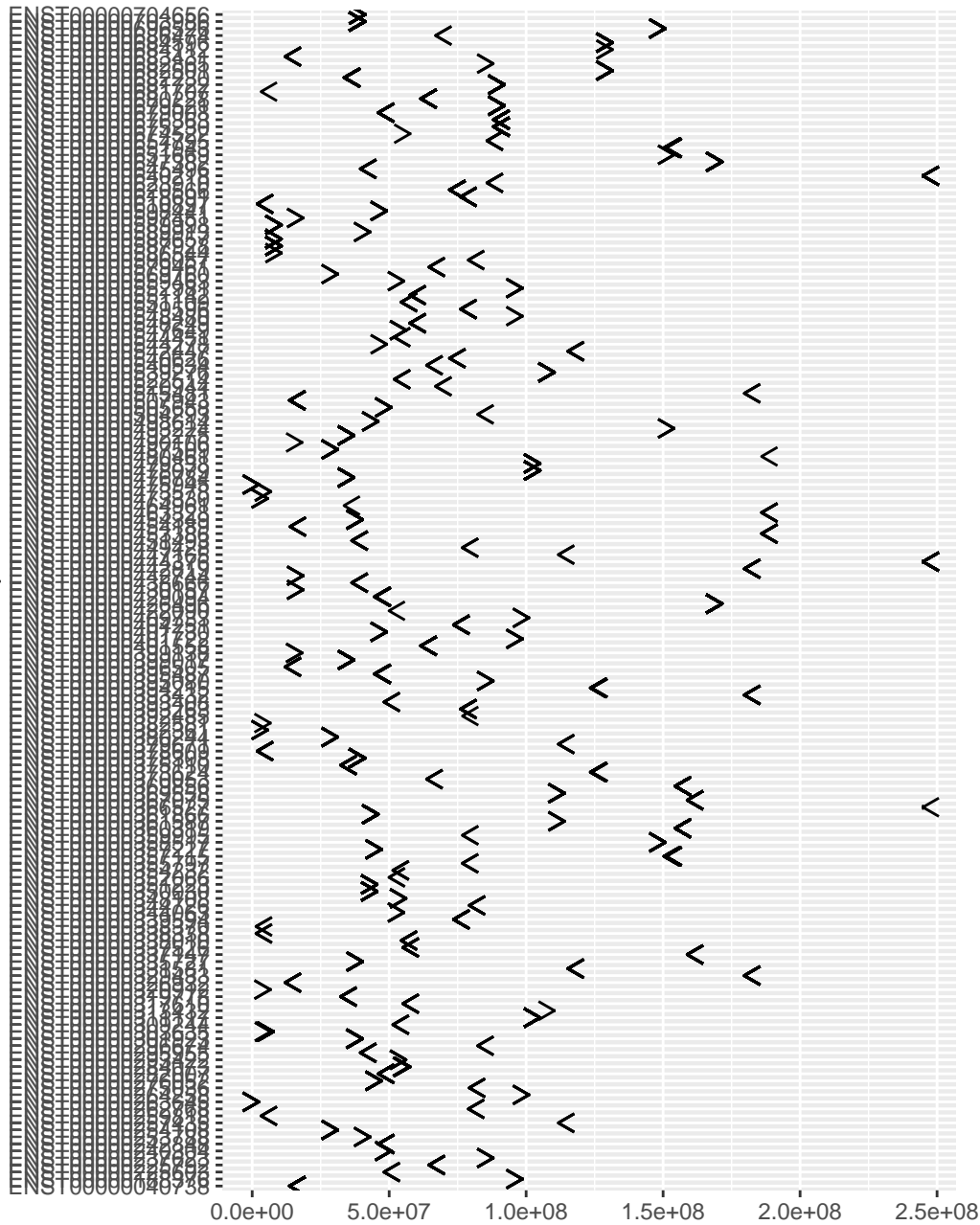
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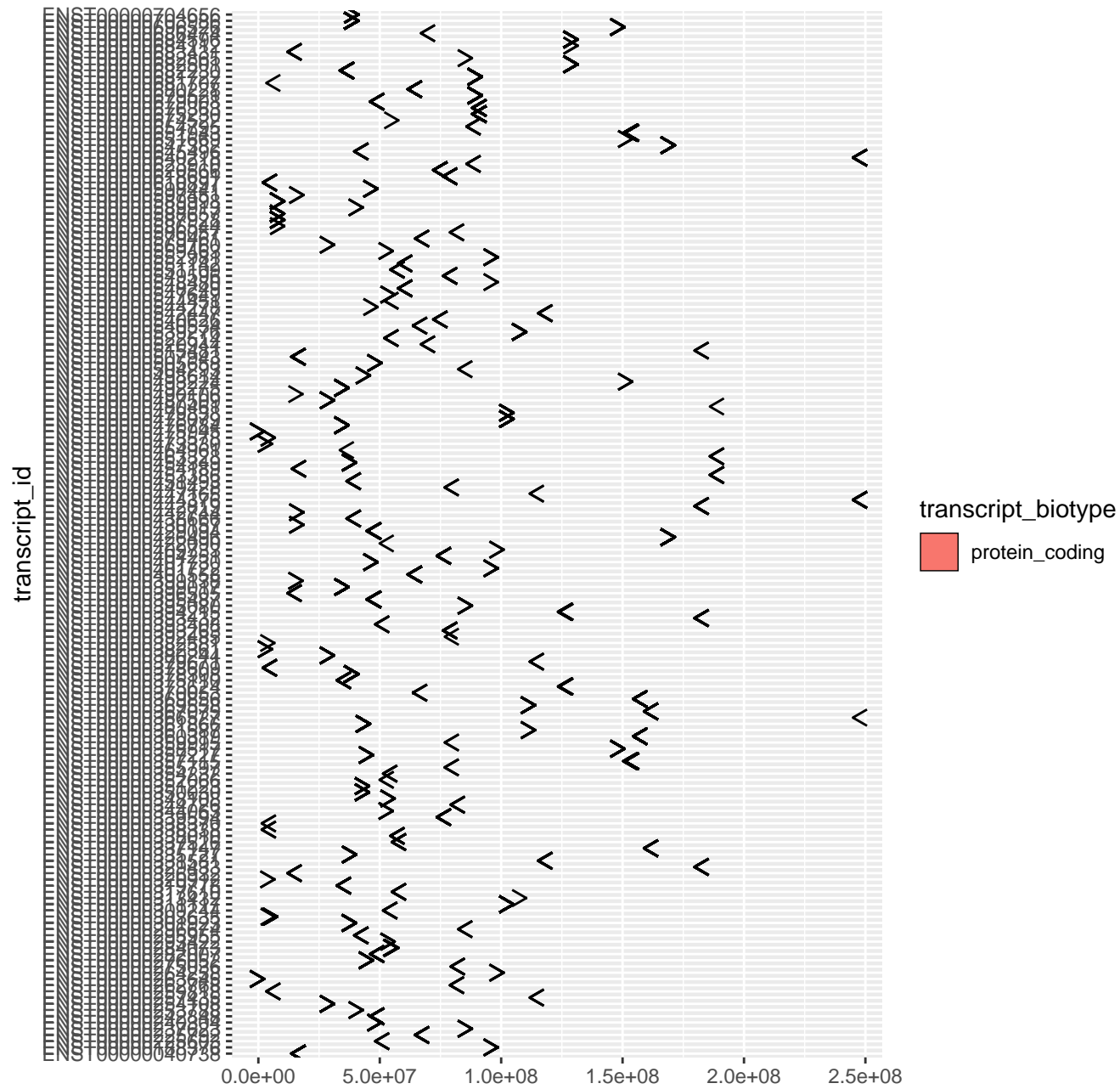


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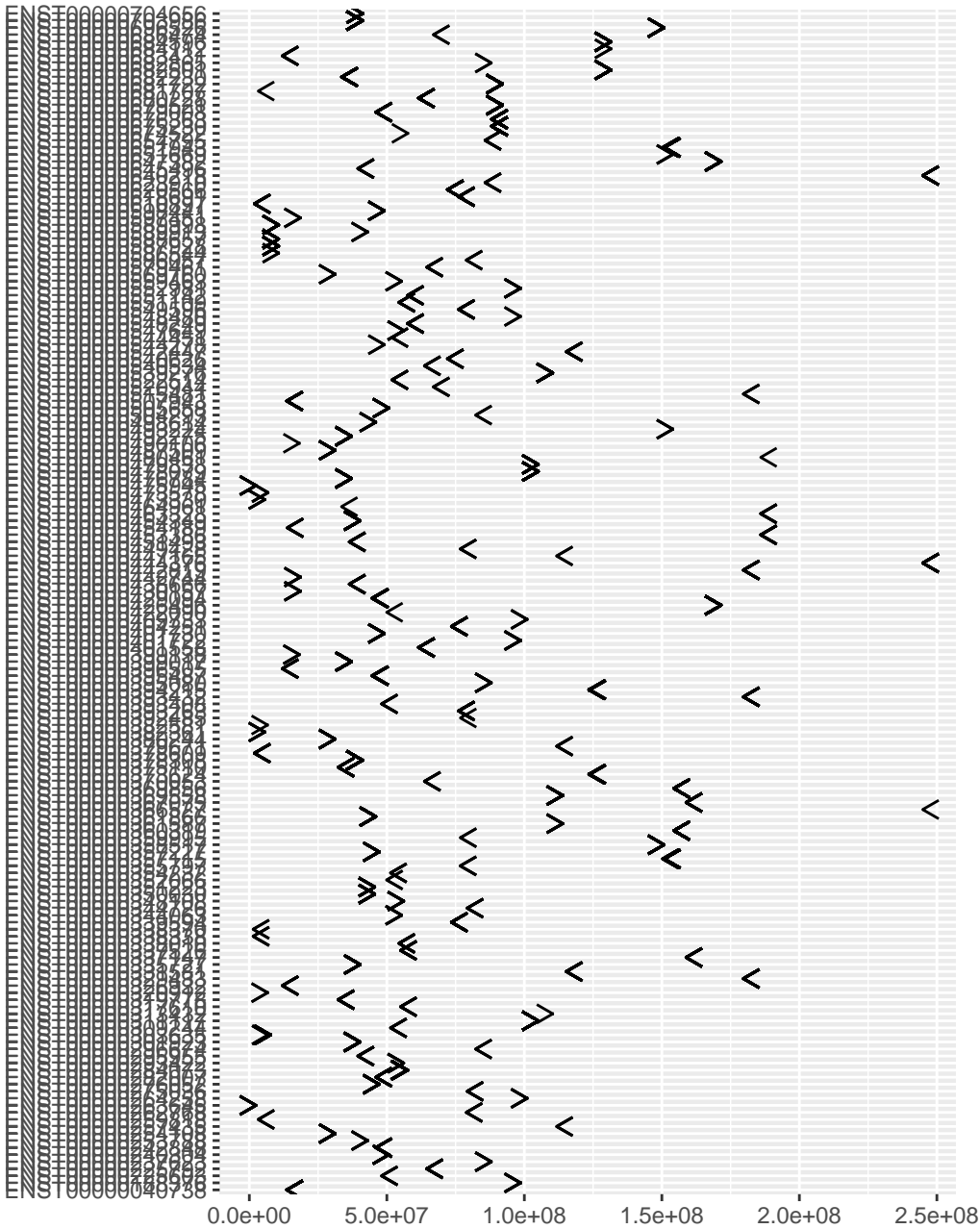


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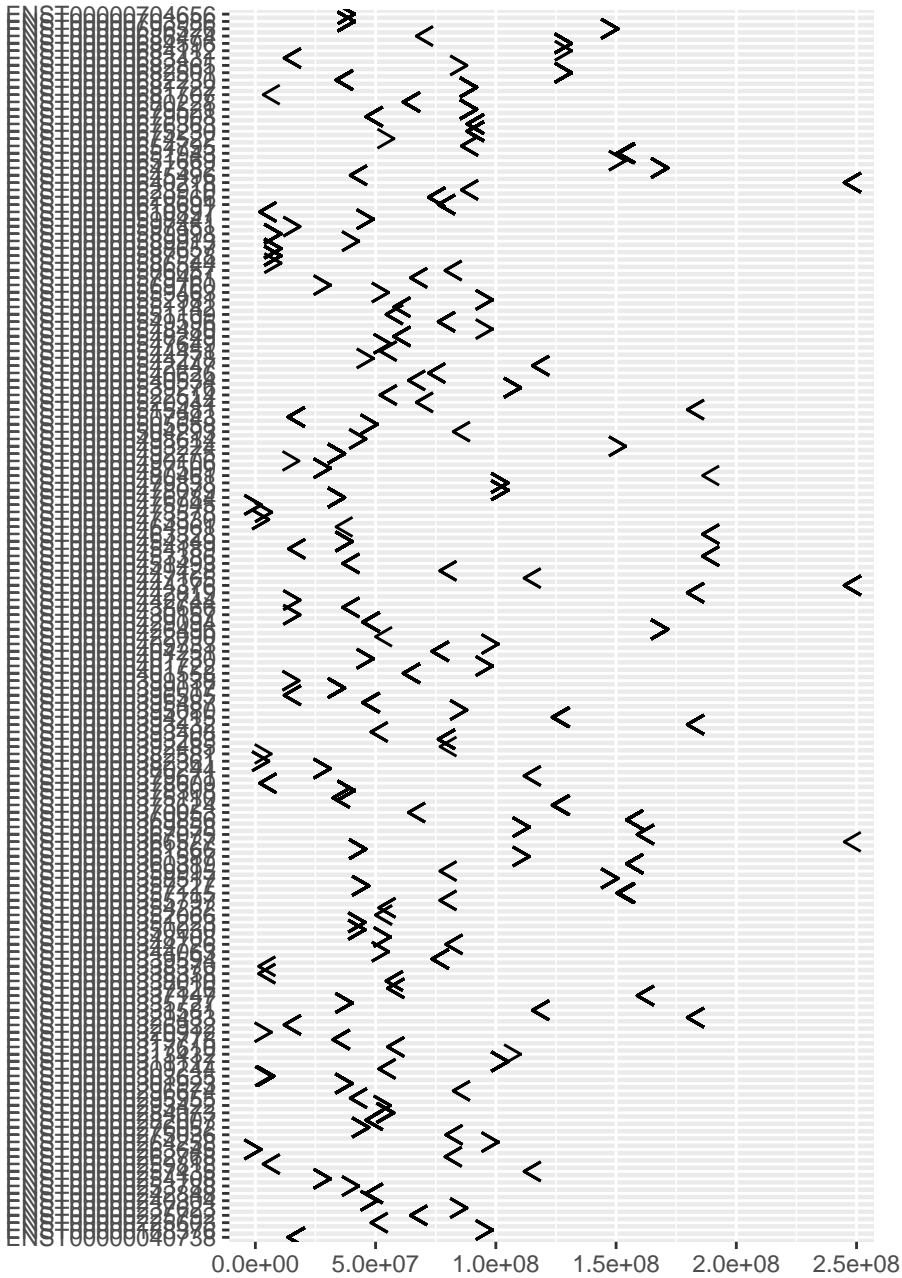
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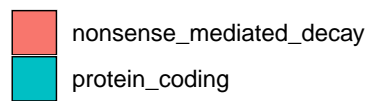


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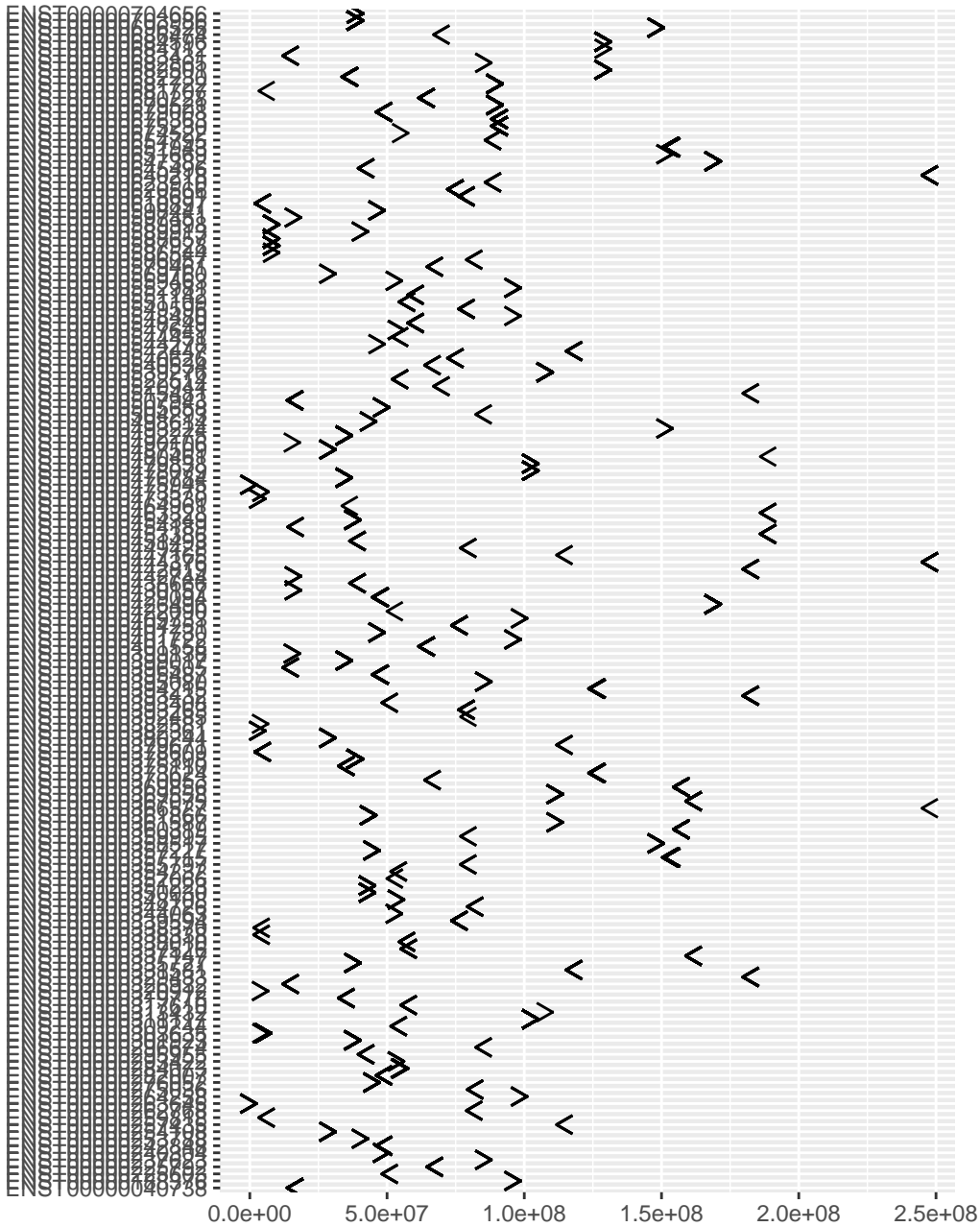


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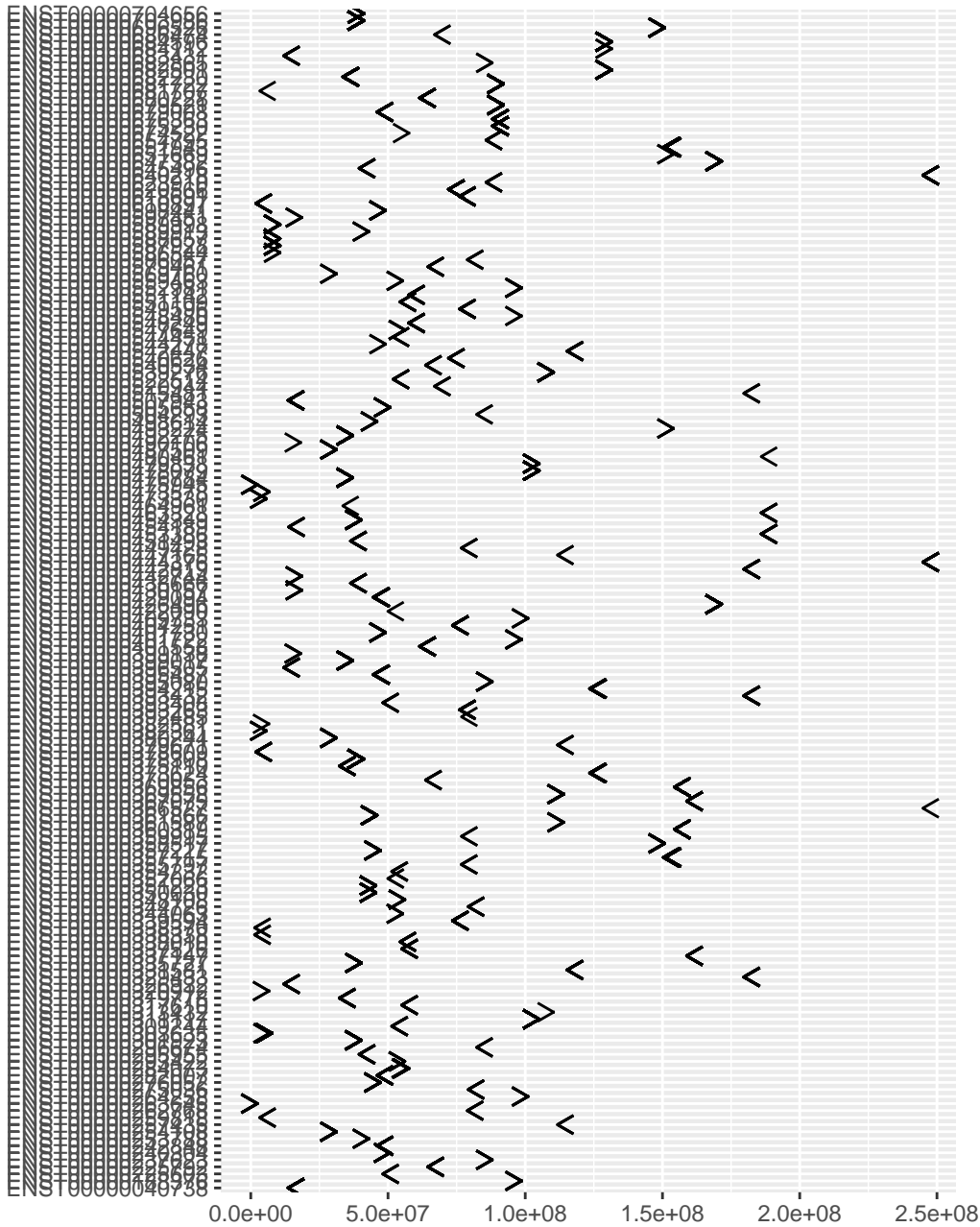


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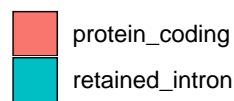
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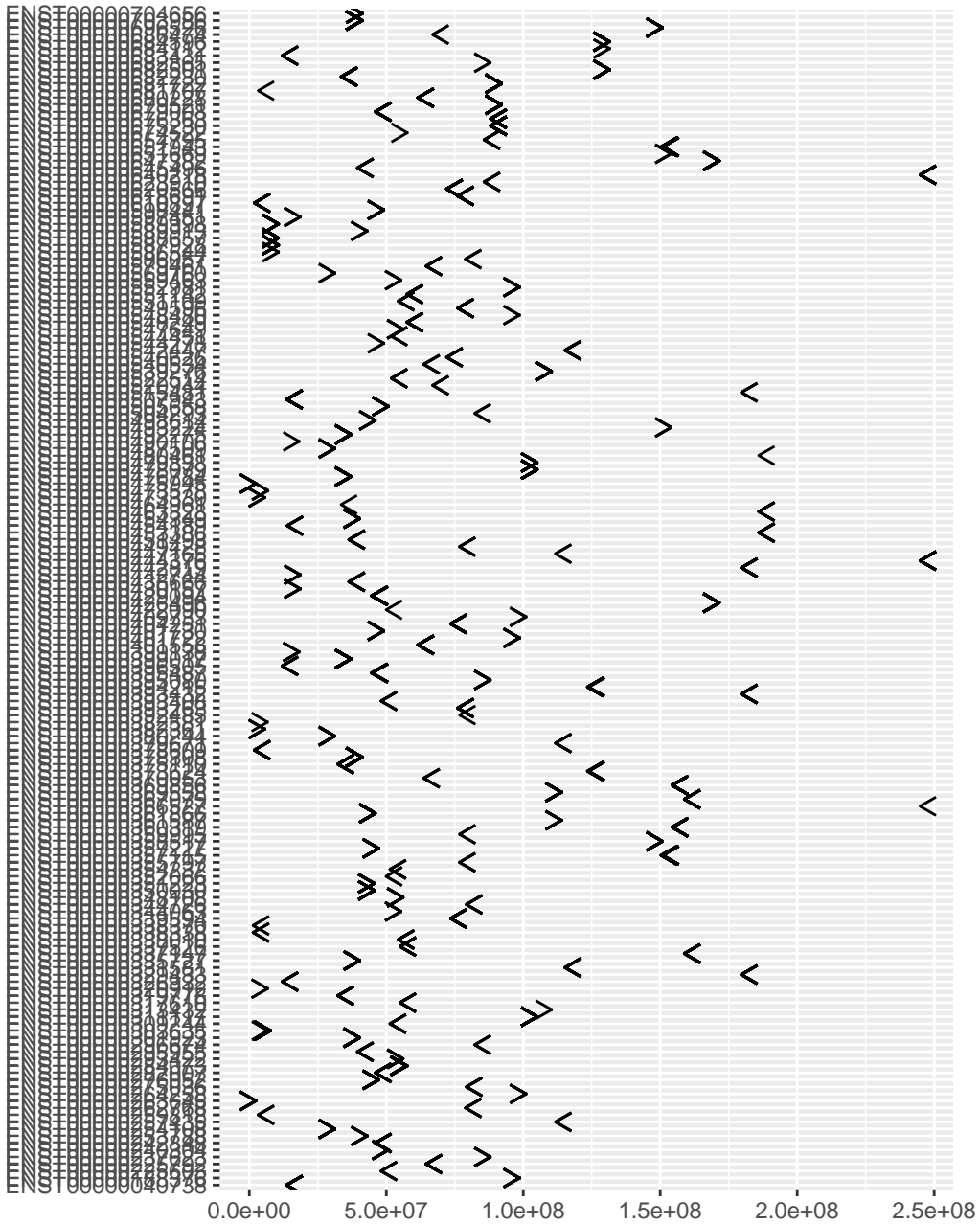


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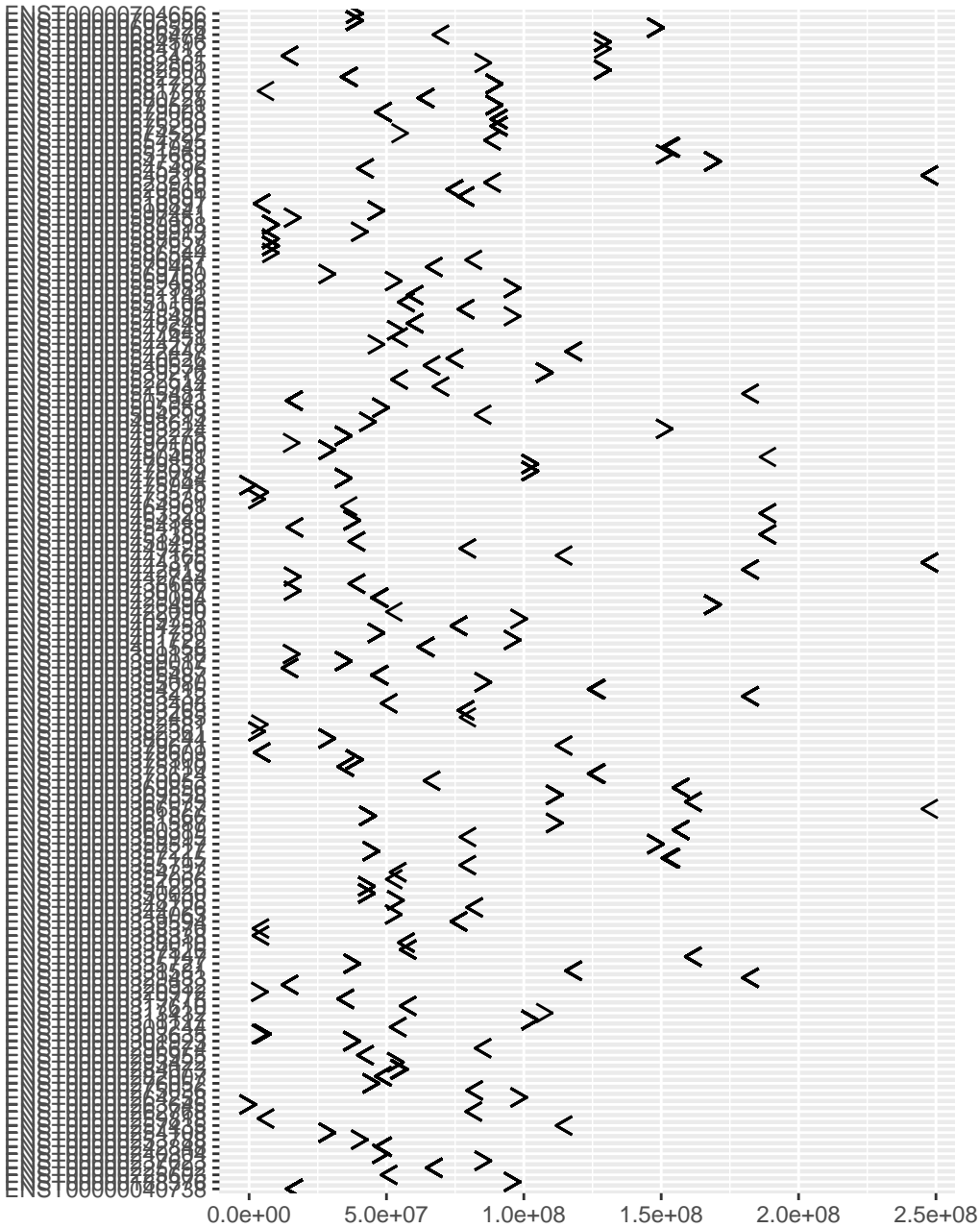
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protein_coding

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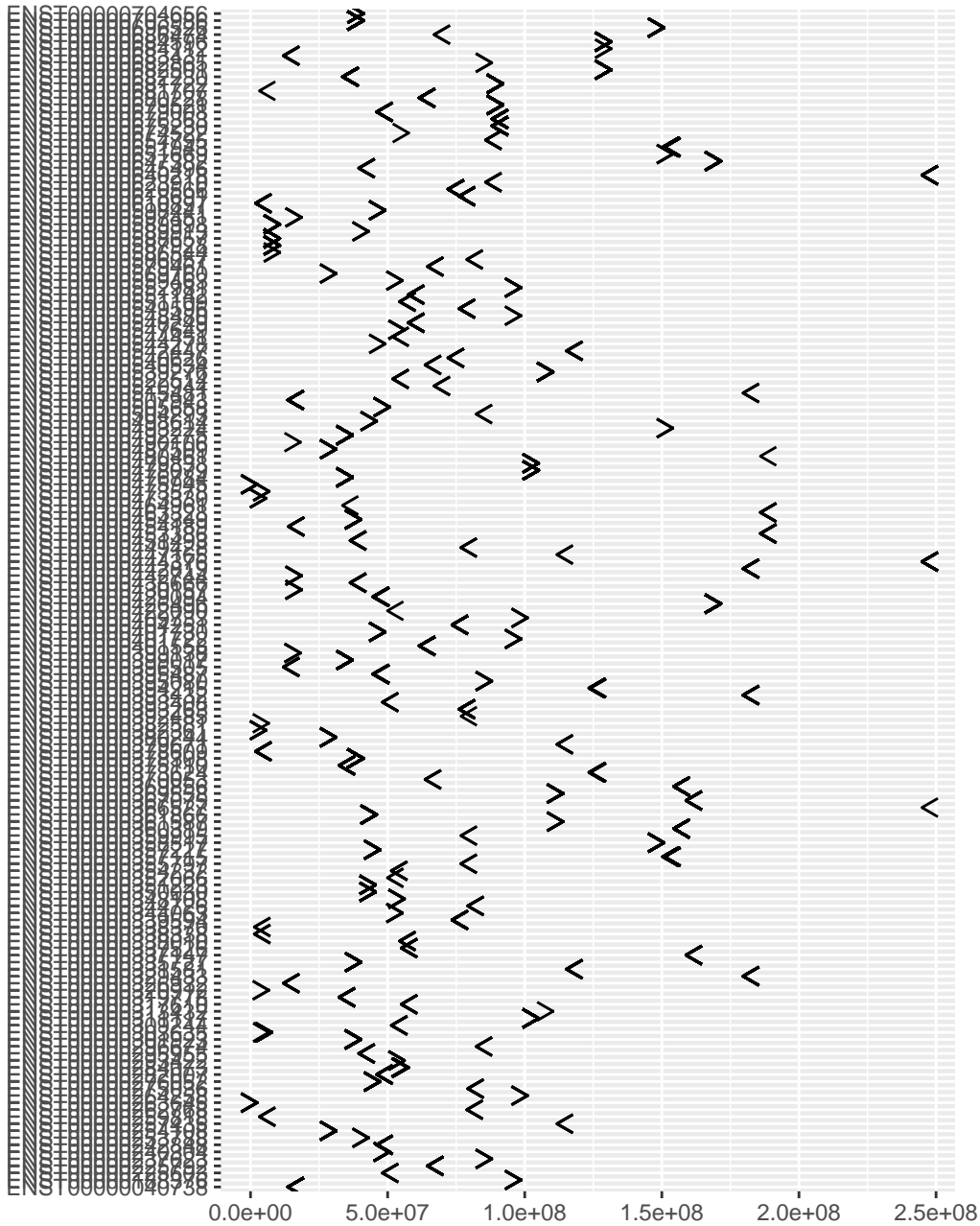


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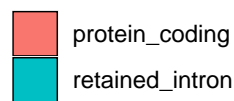
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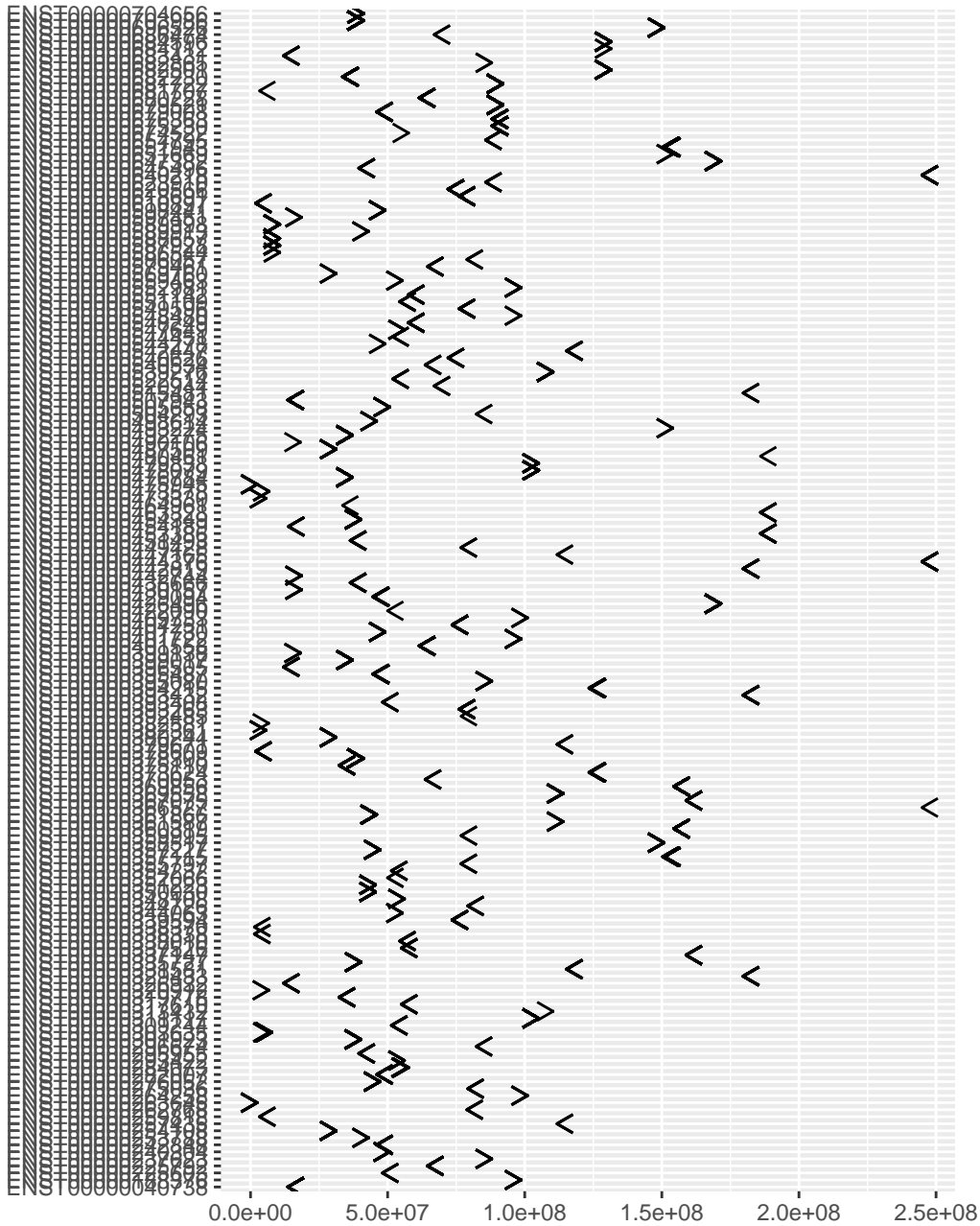


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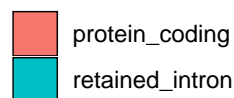


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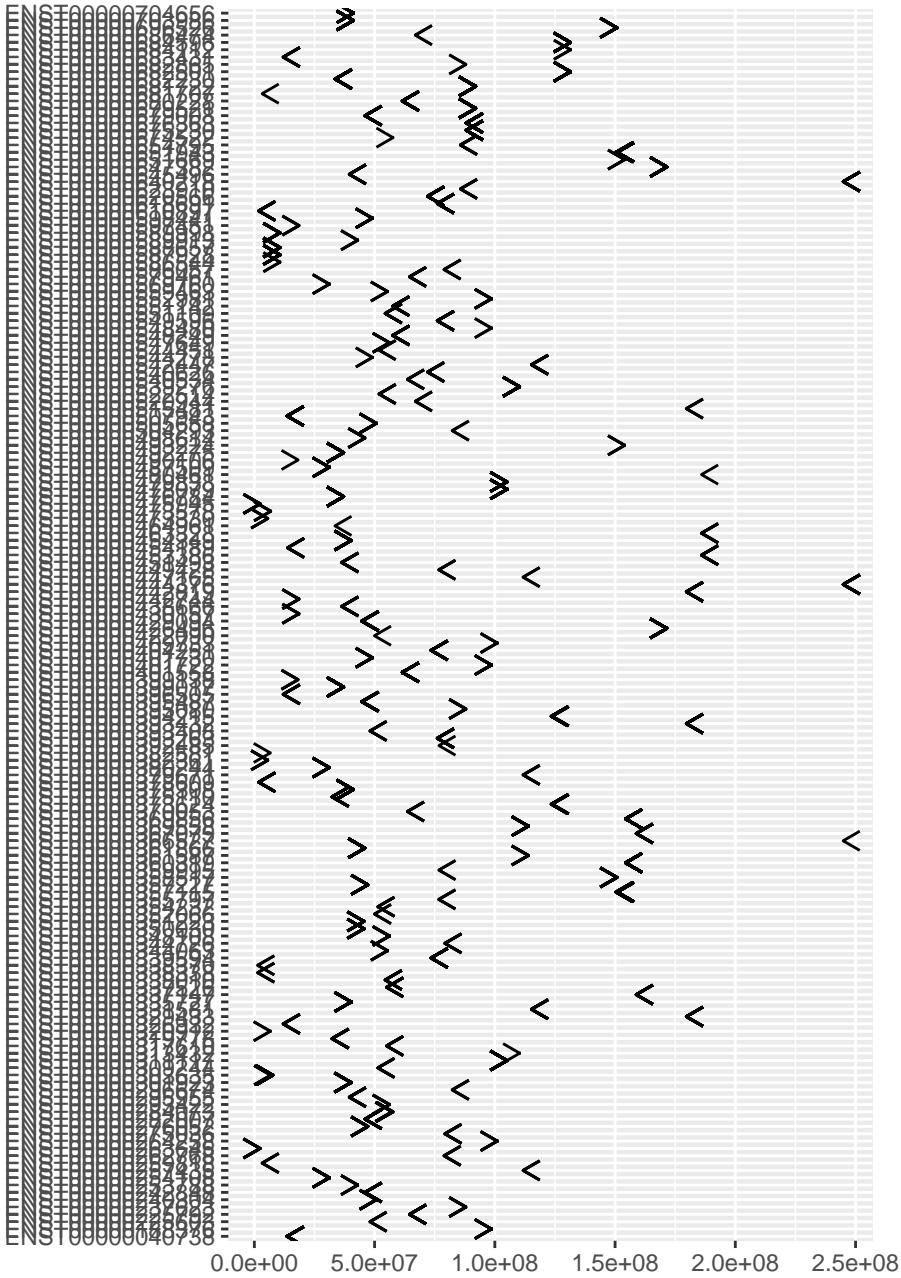


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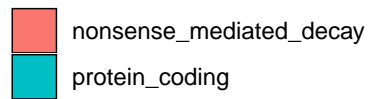


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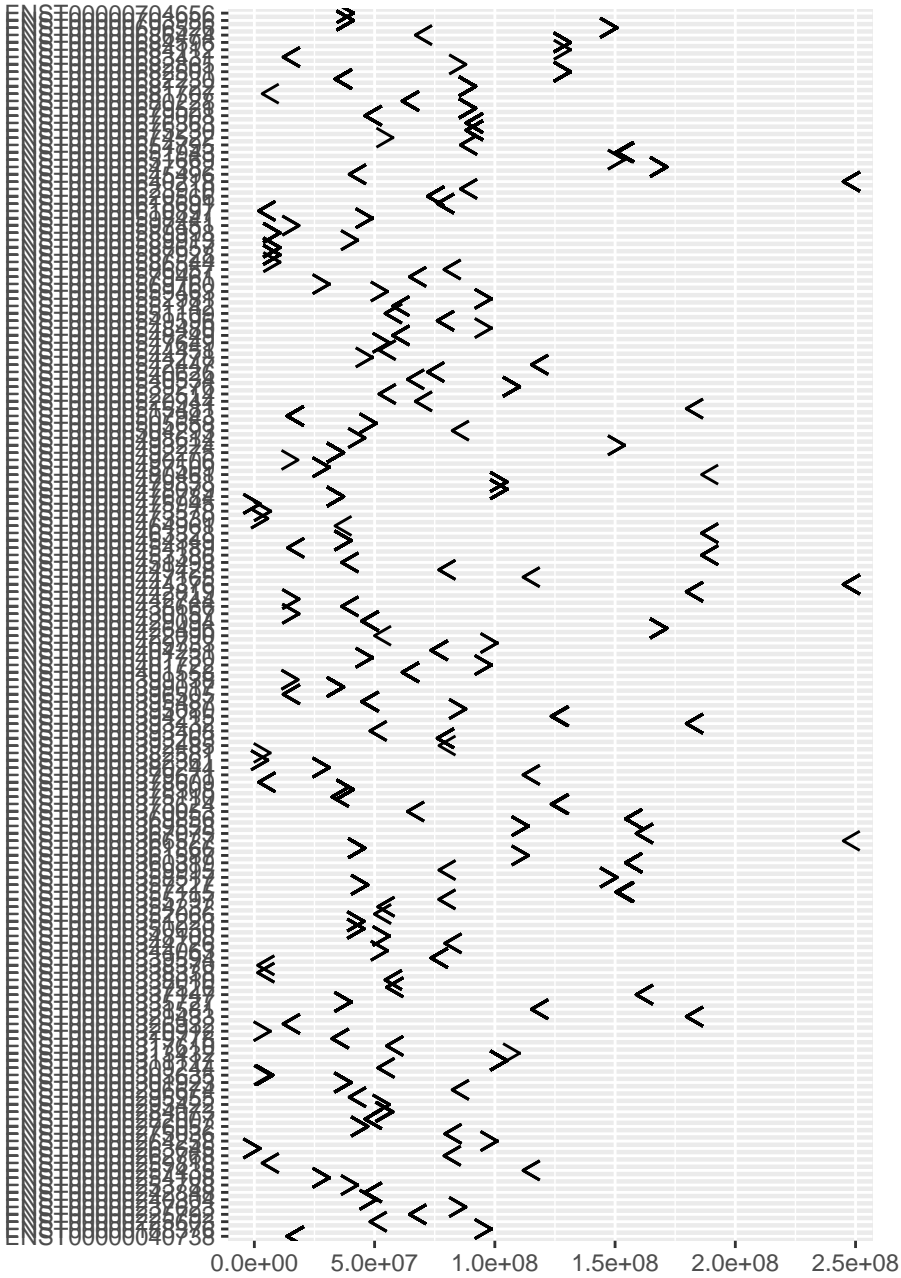


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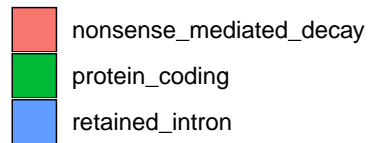


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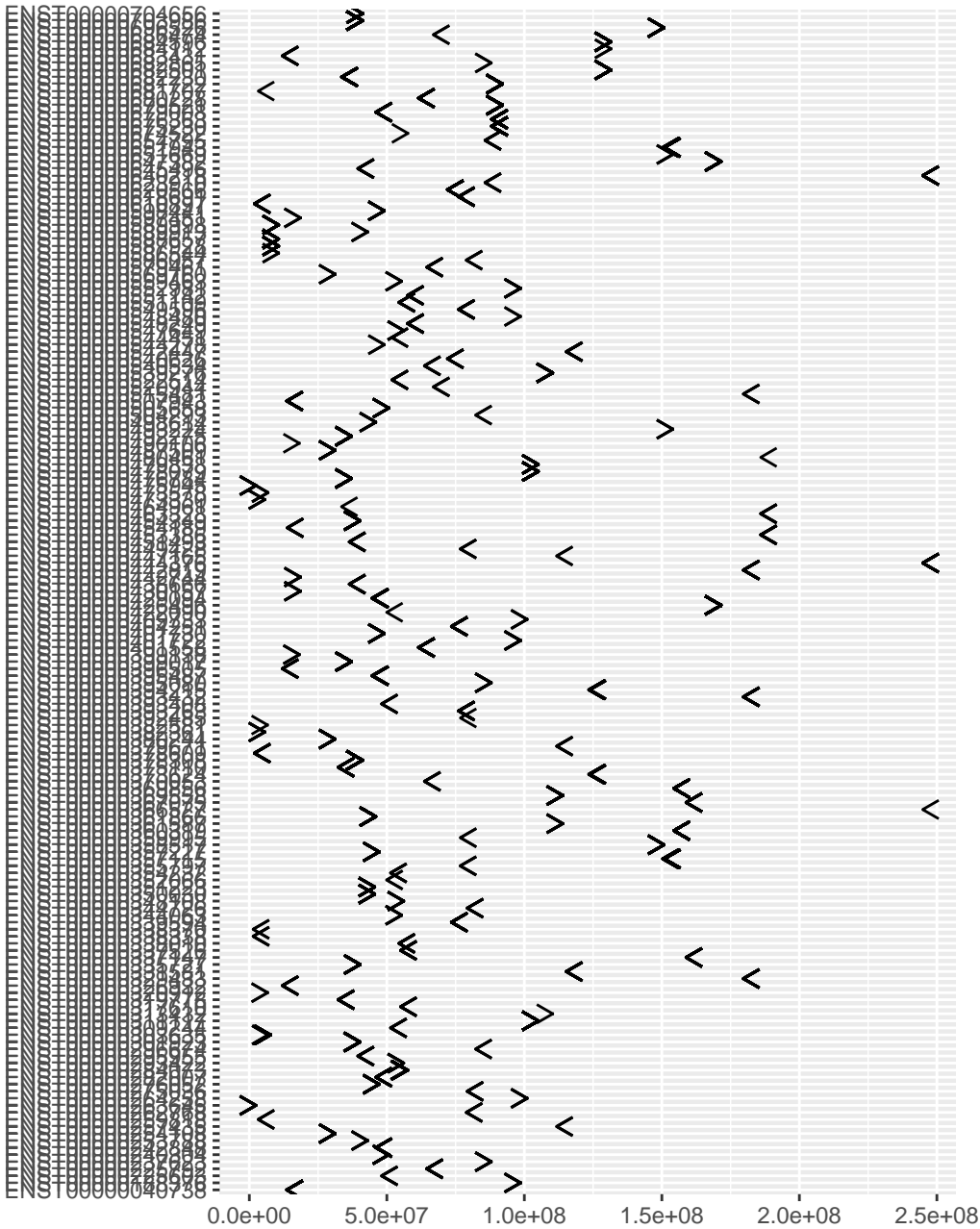


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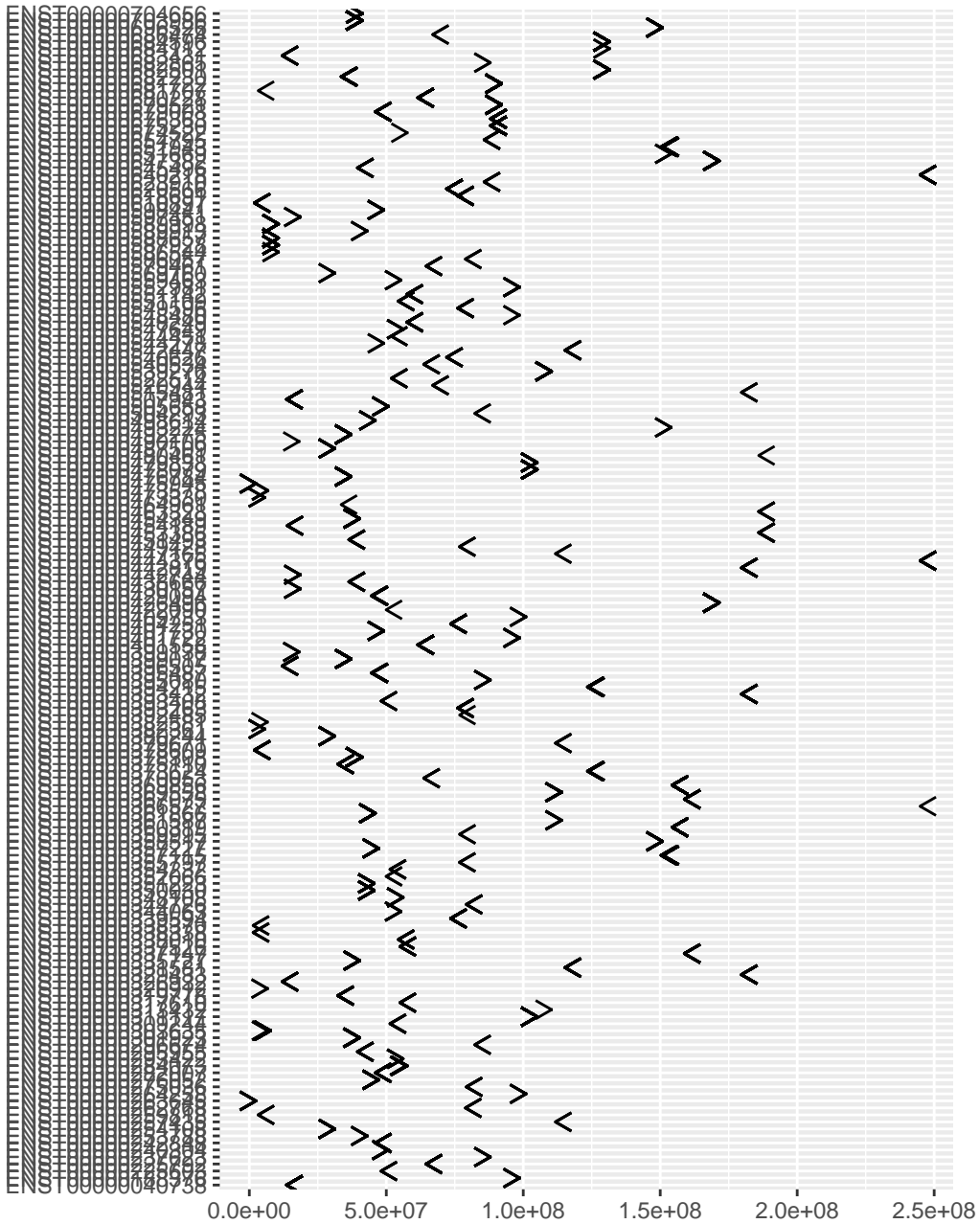


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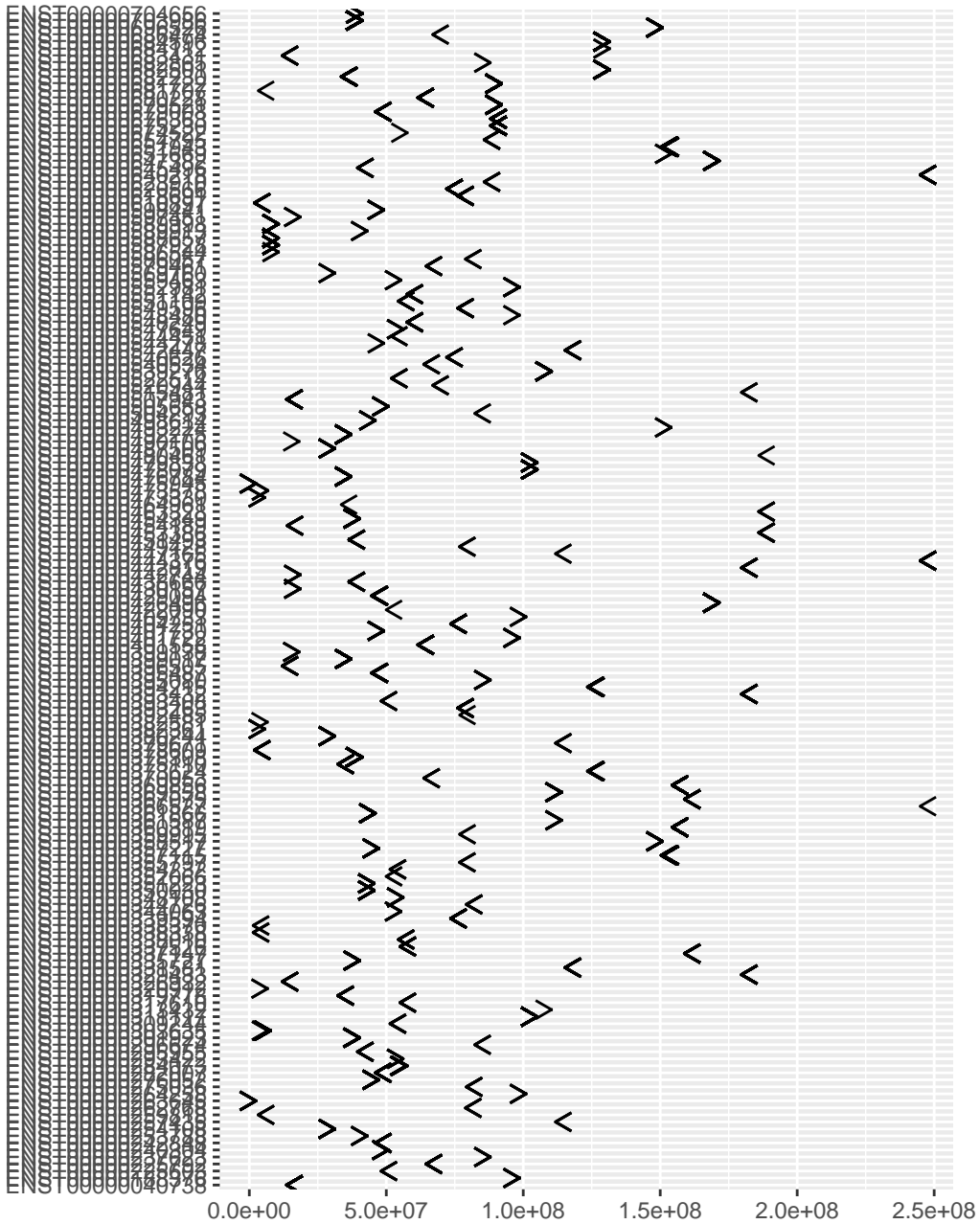
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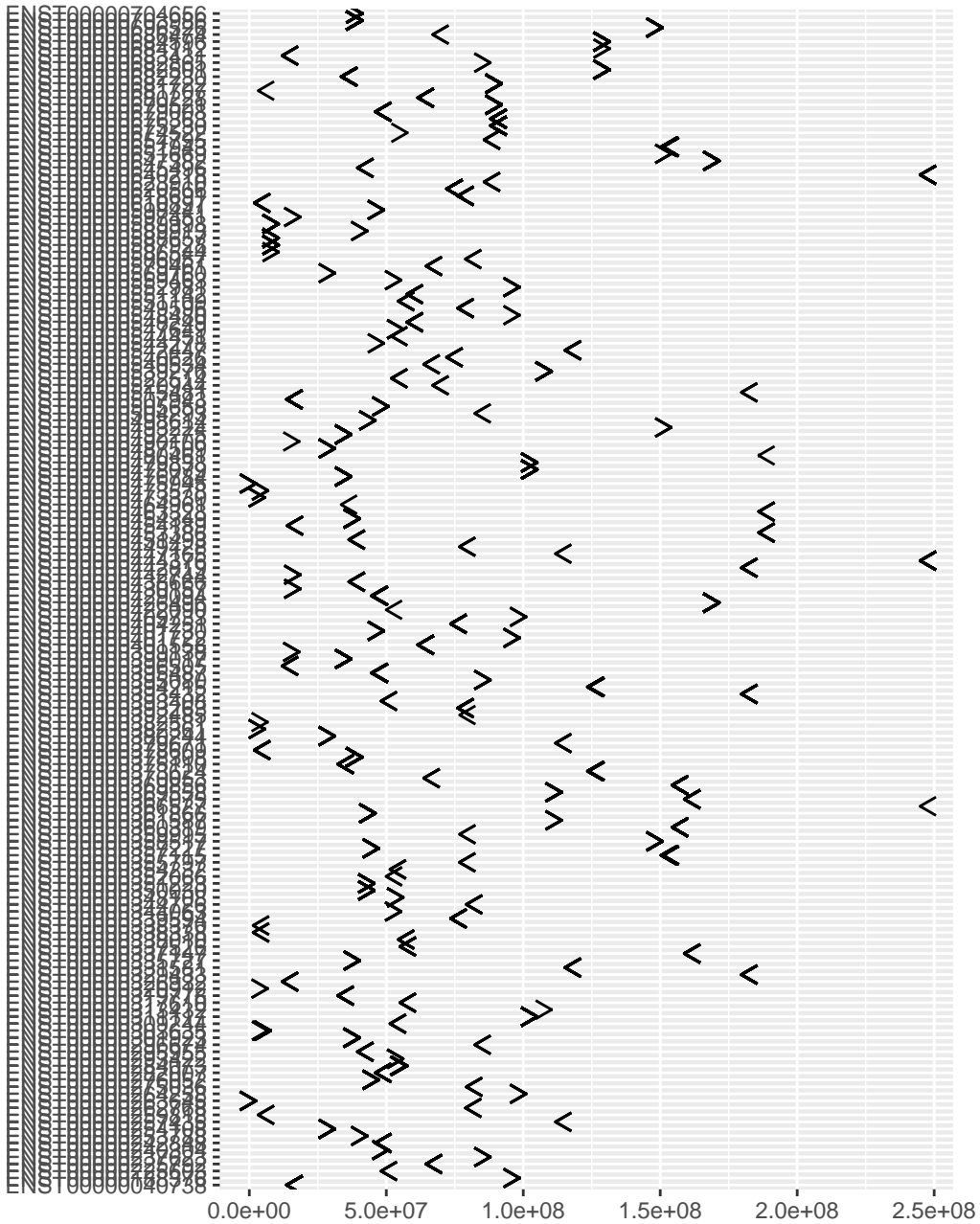
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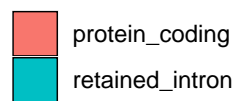
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TRA2B

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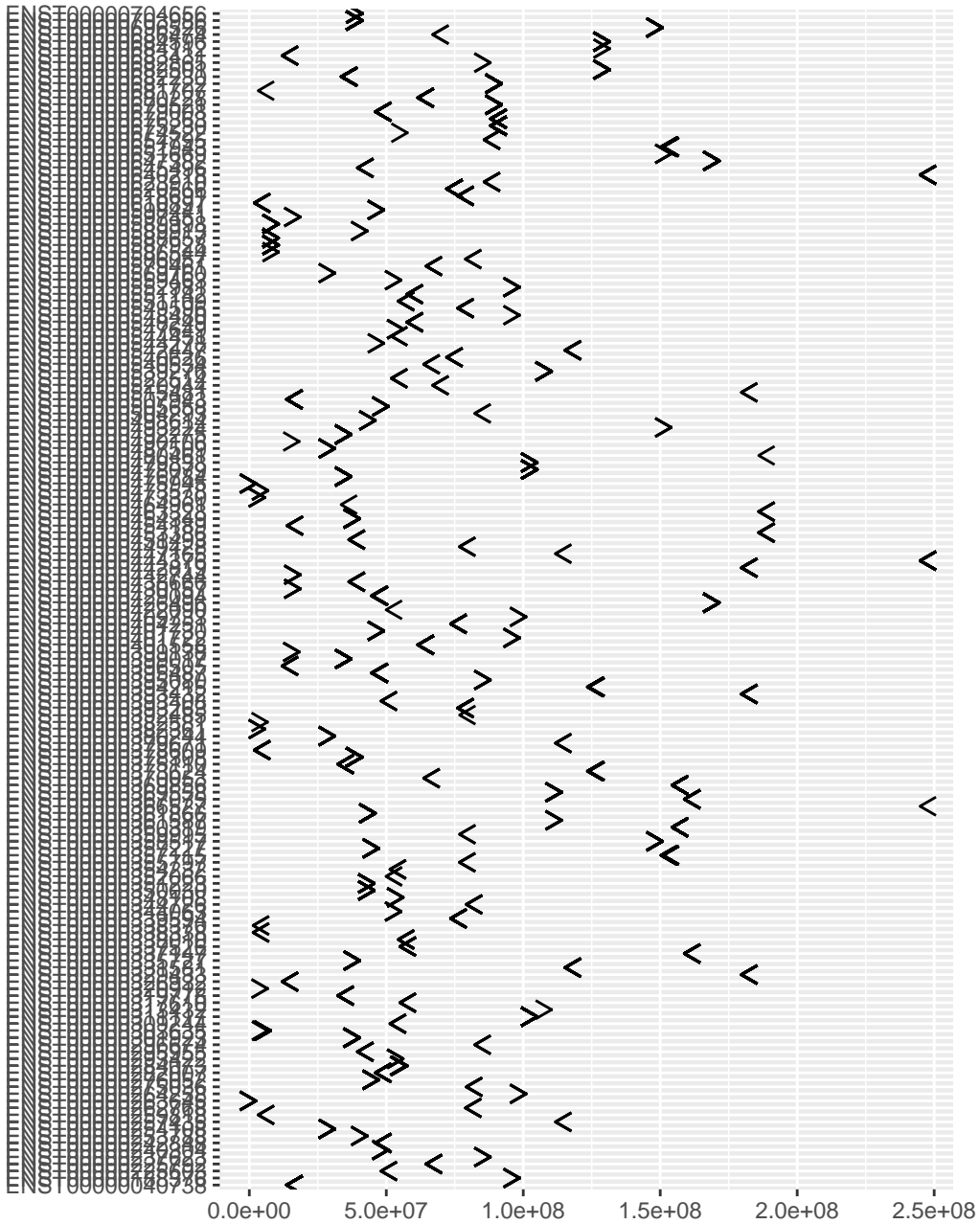


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GPM6B

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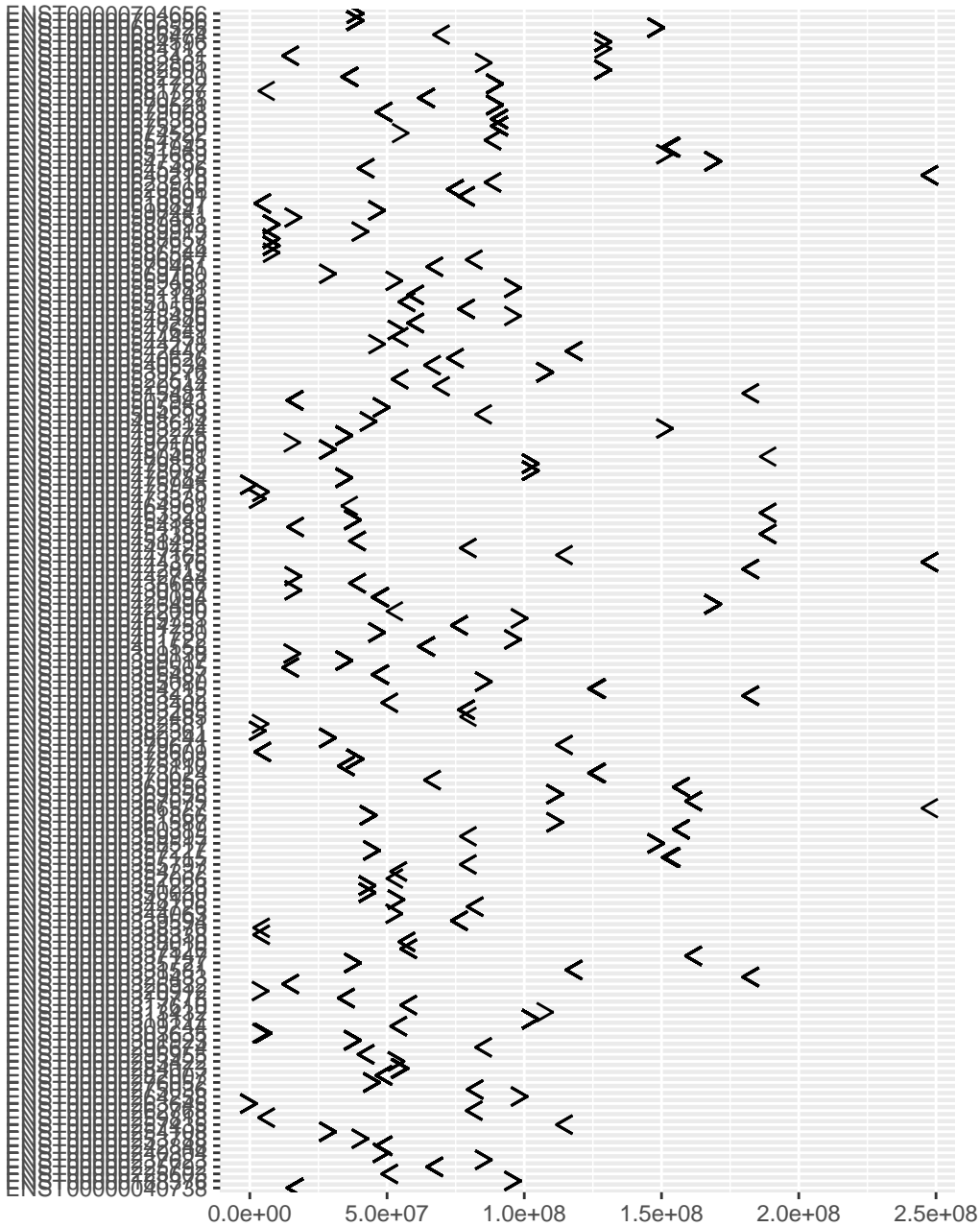


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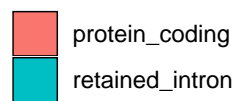
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CLASP2

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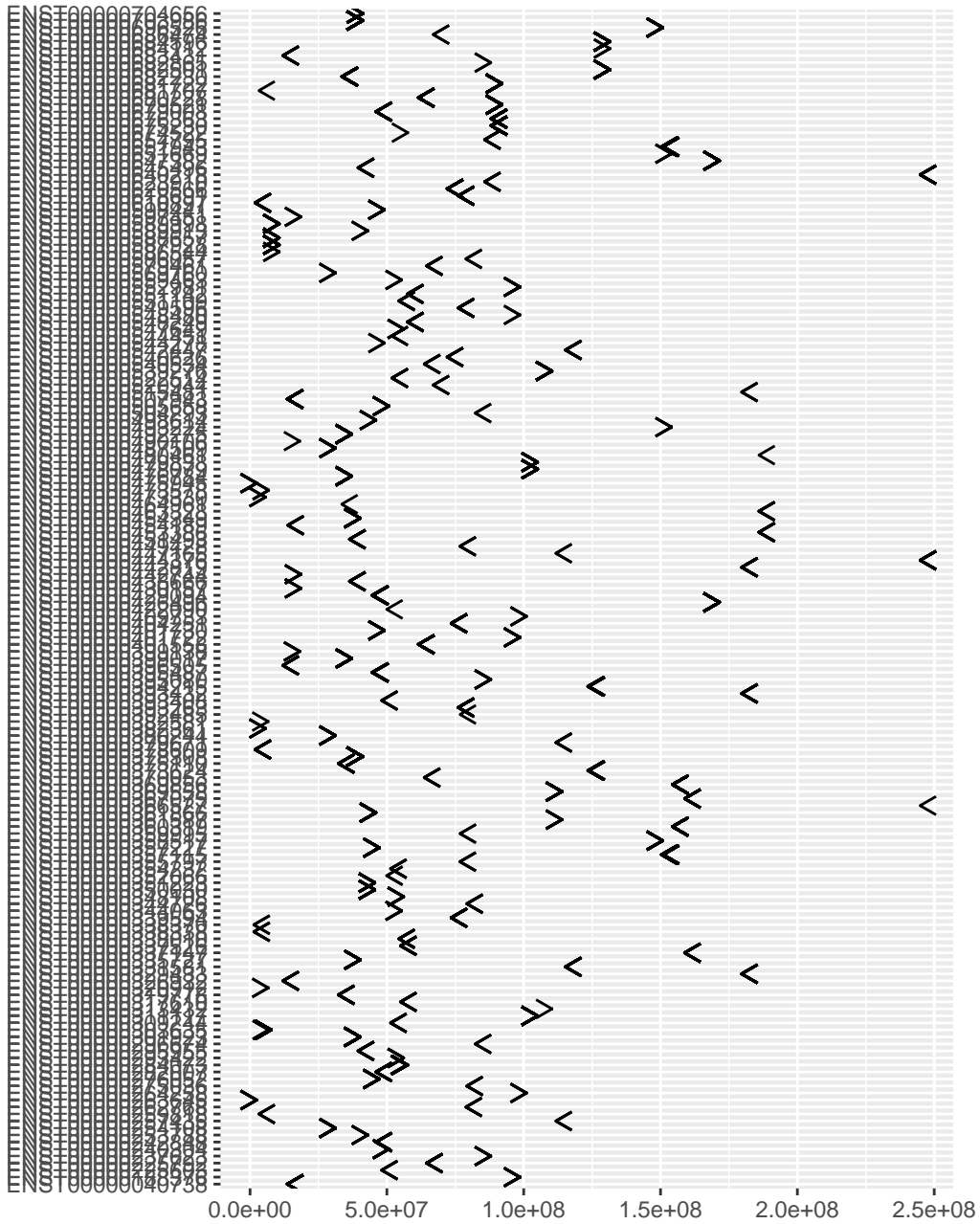


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RPS27

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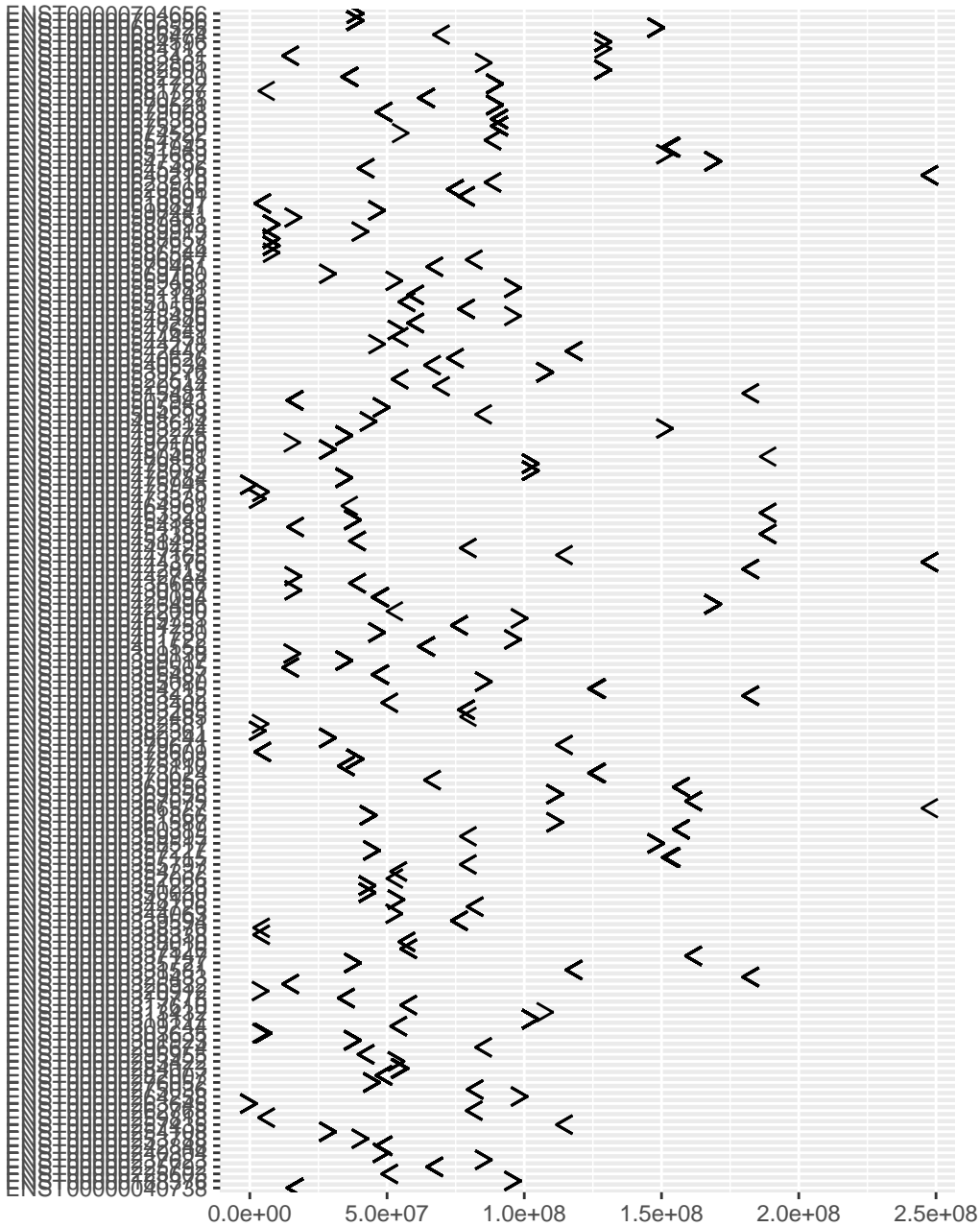
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 protein_coding

retained_intron

SNHG6

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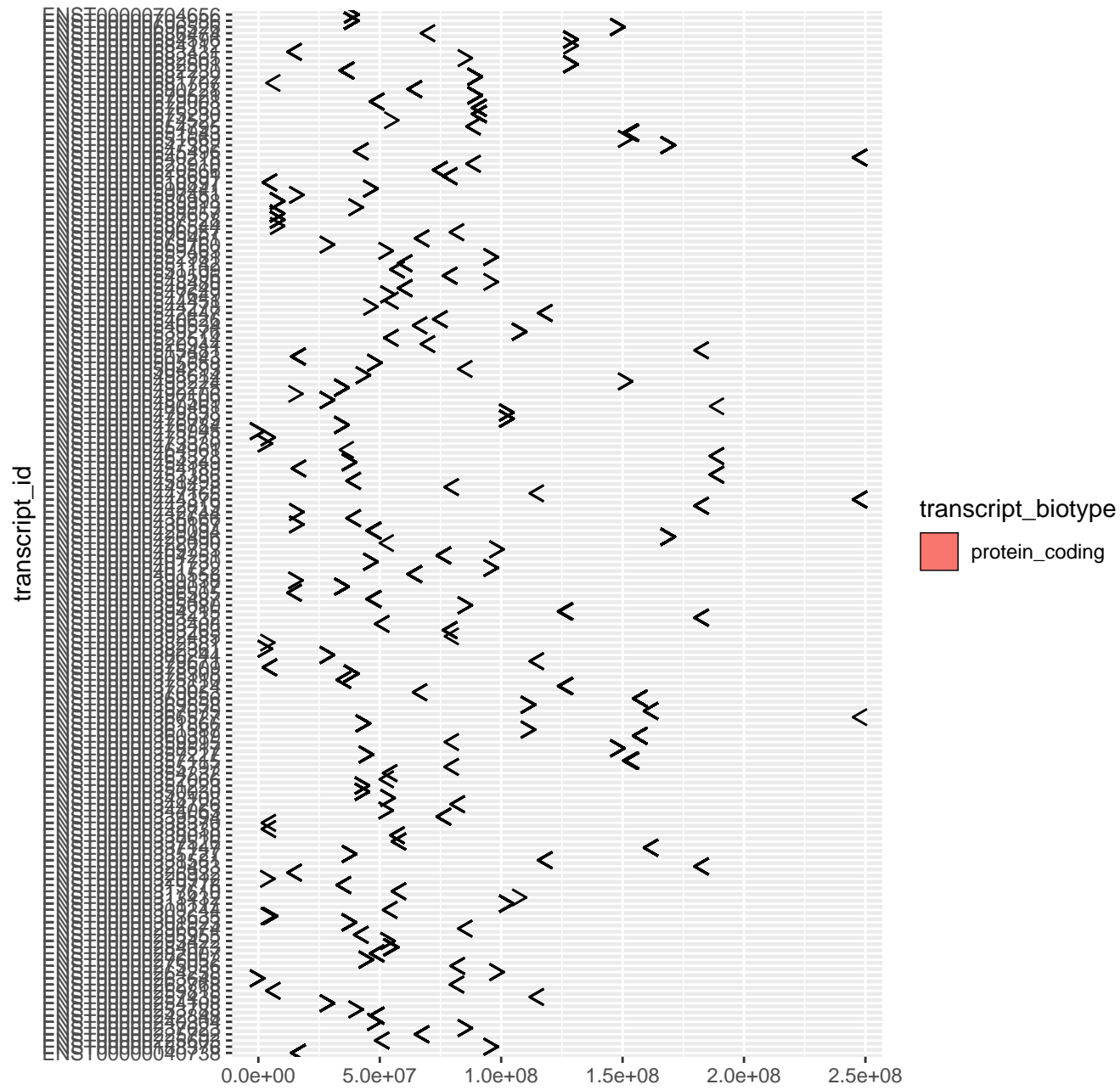


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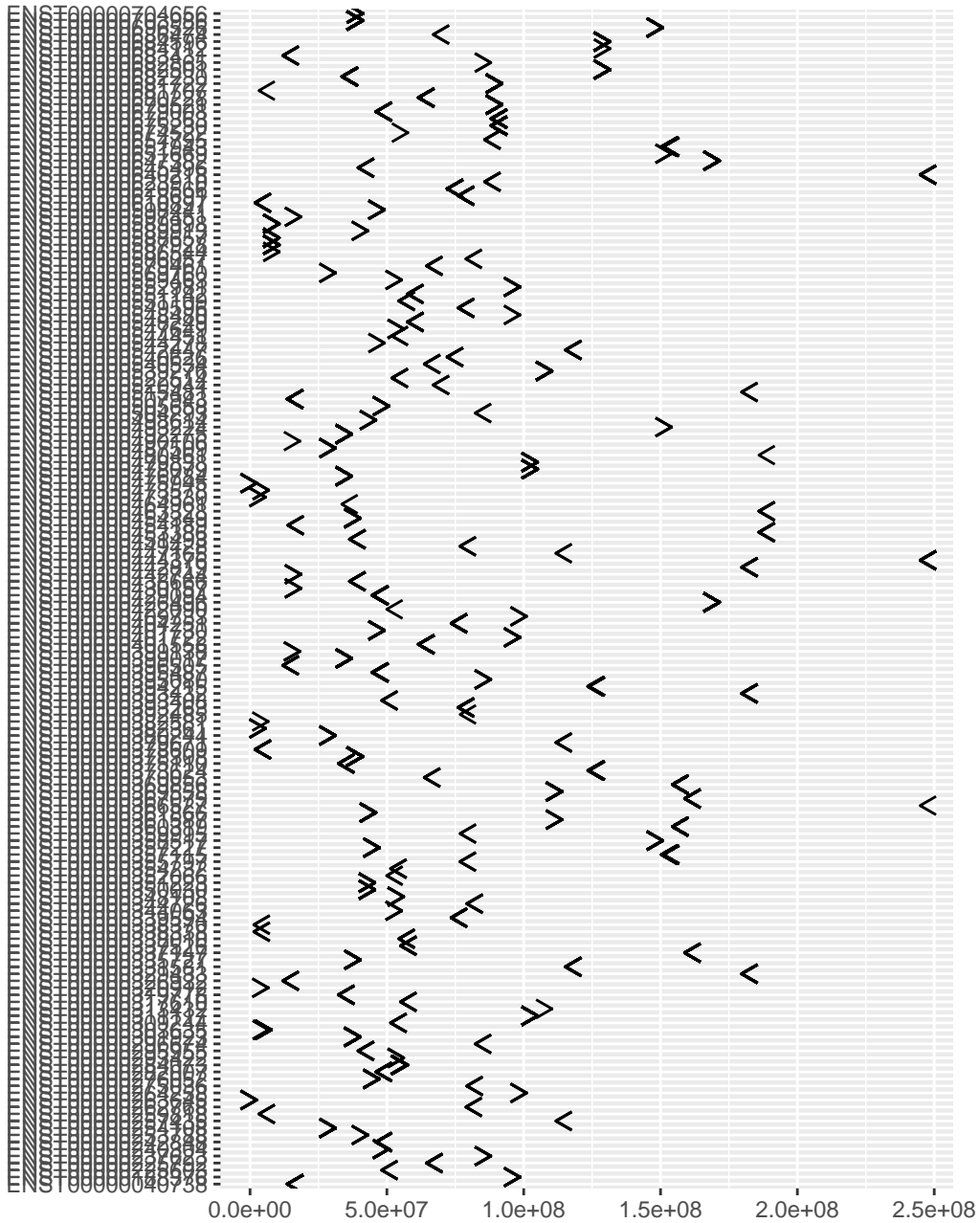
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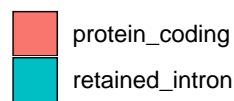


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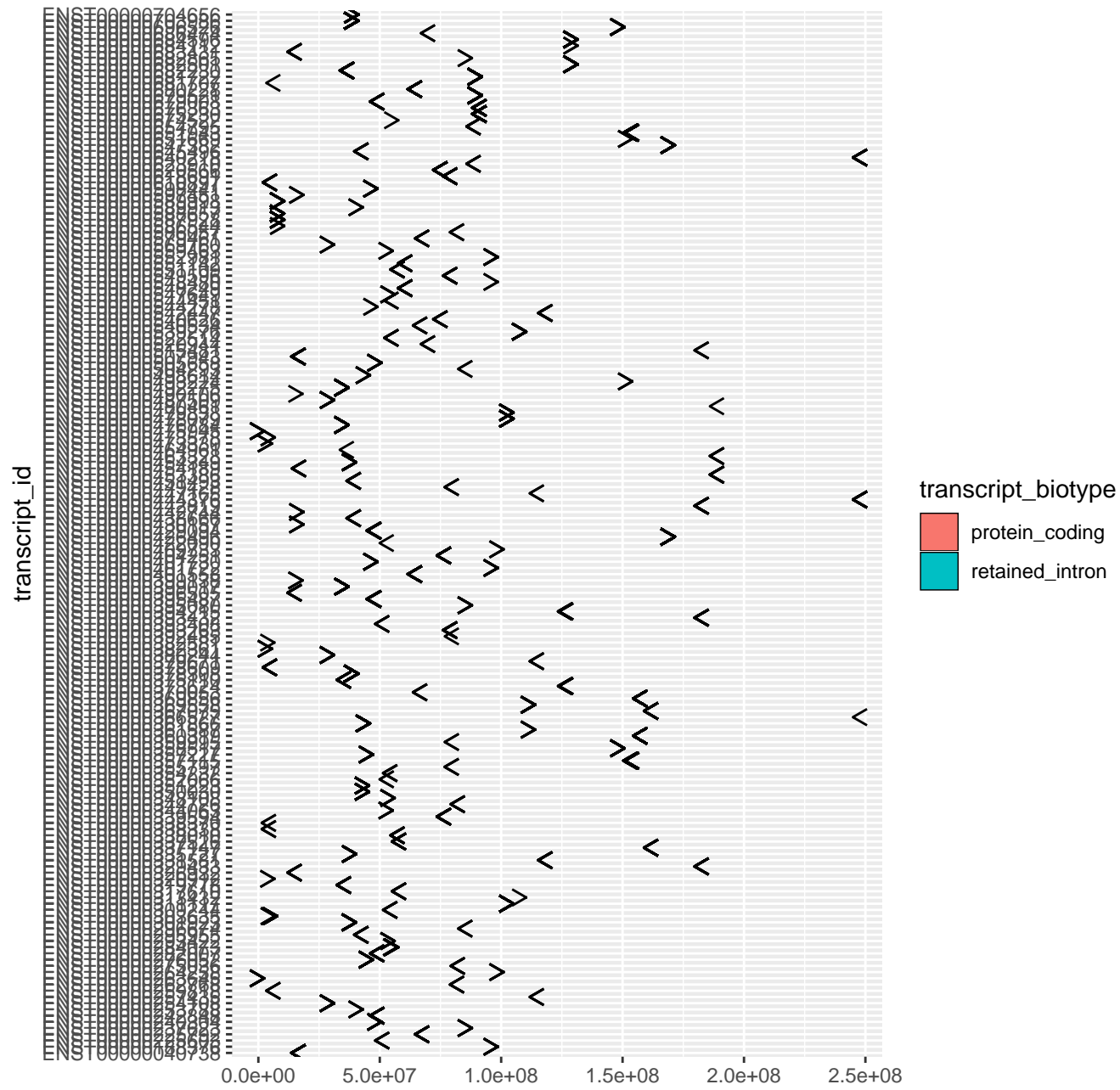
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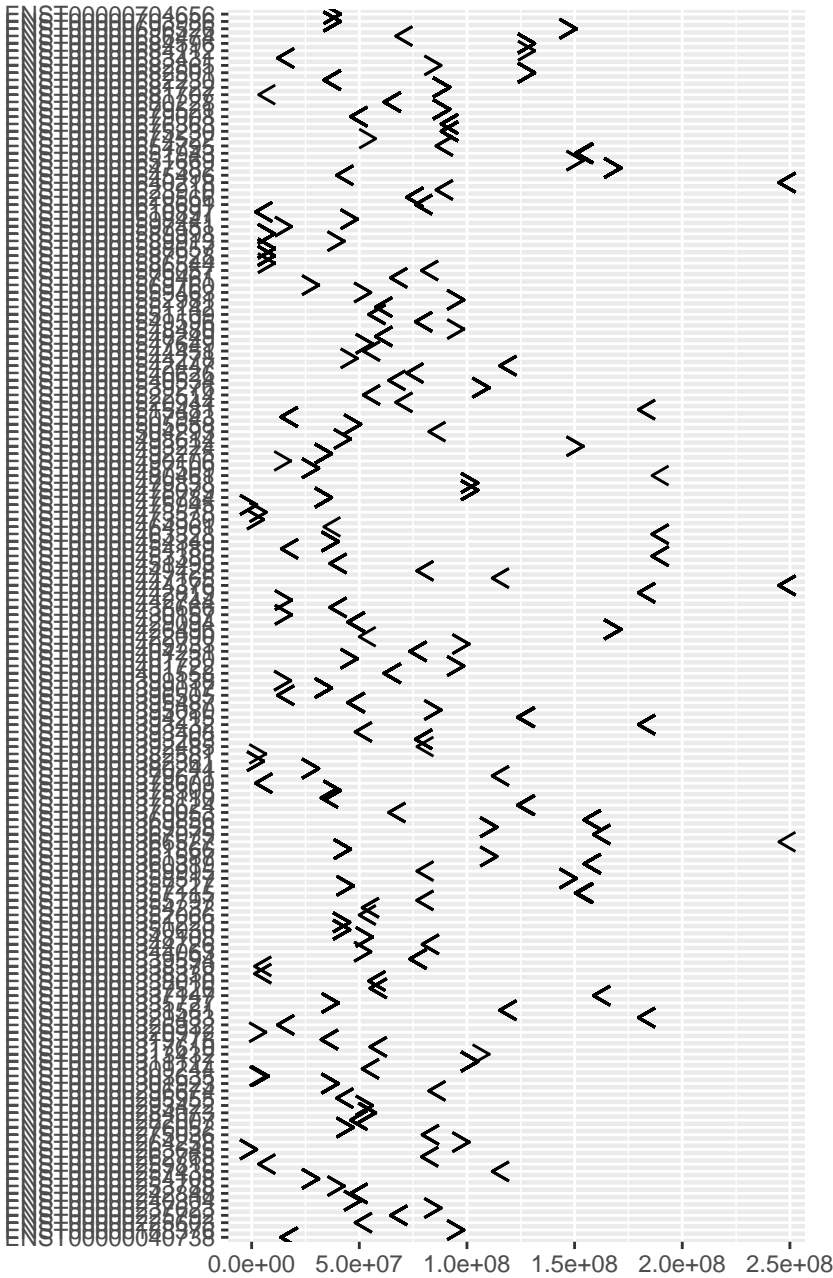


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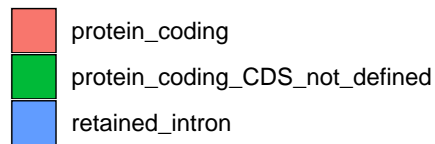


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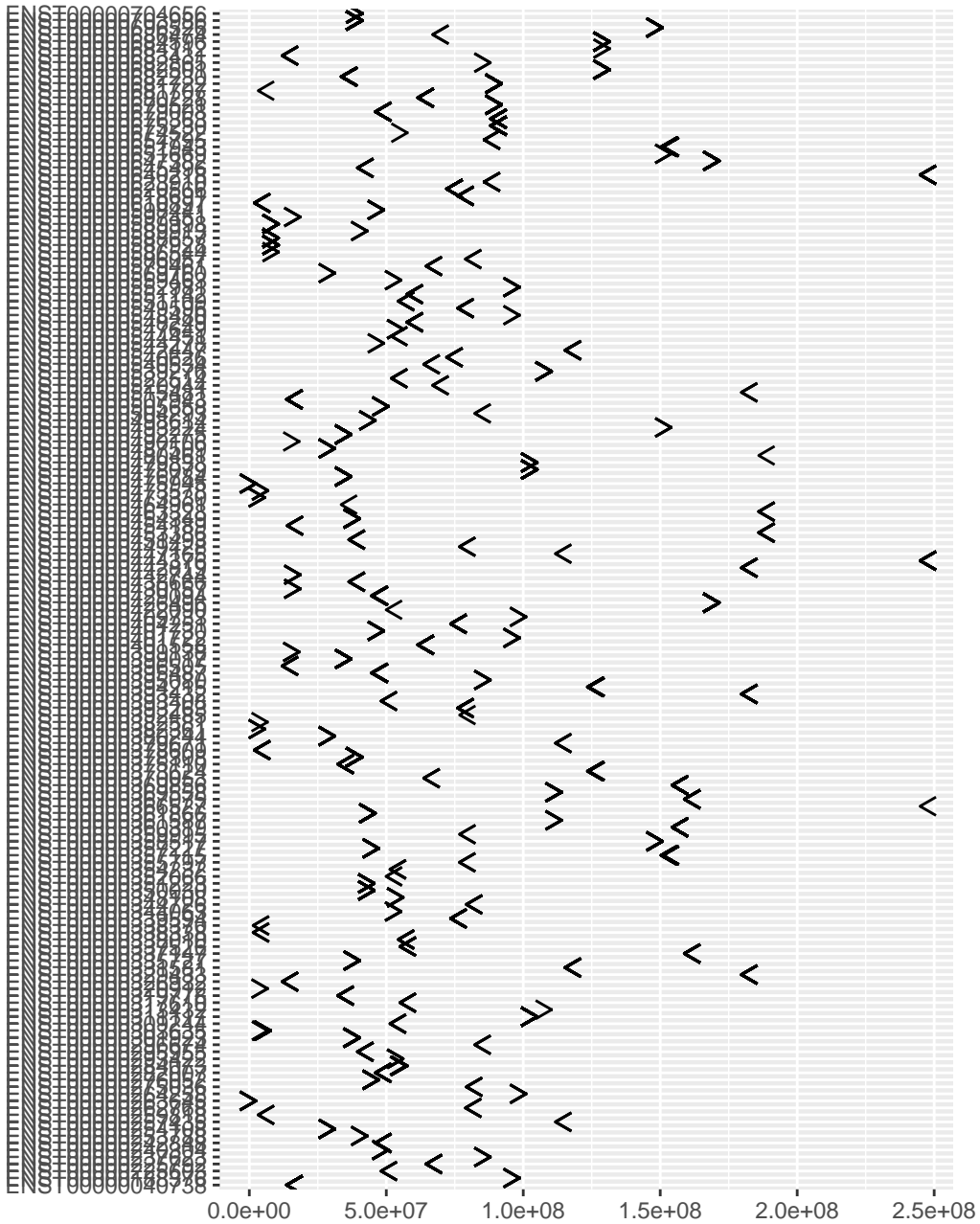


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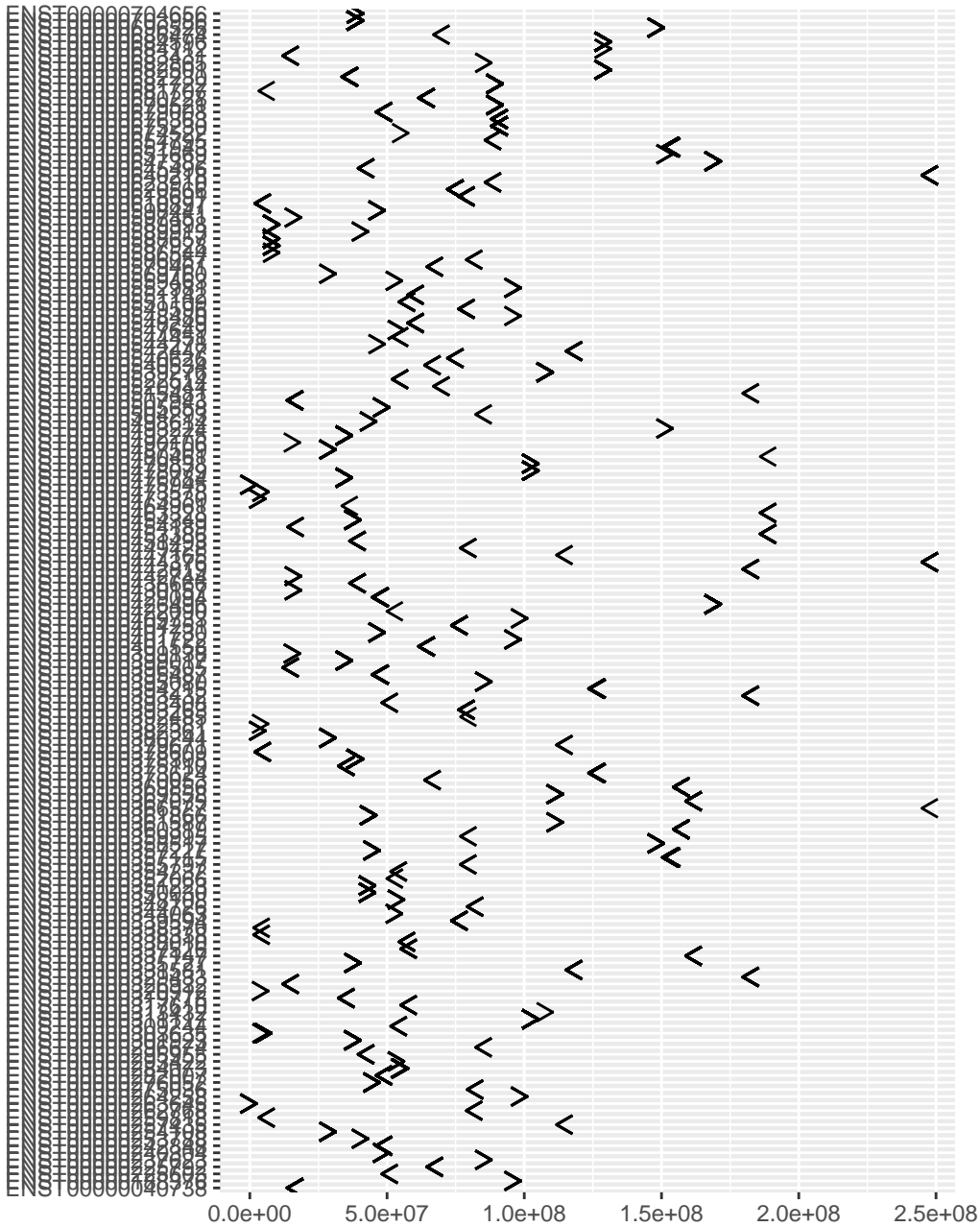
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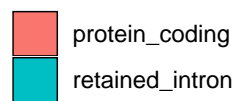
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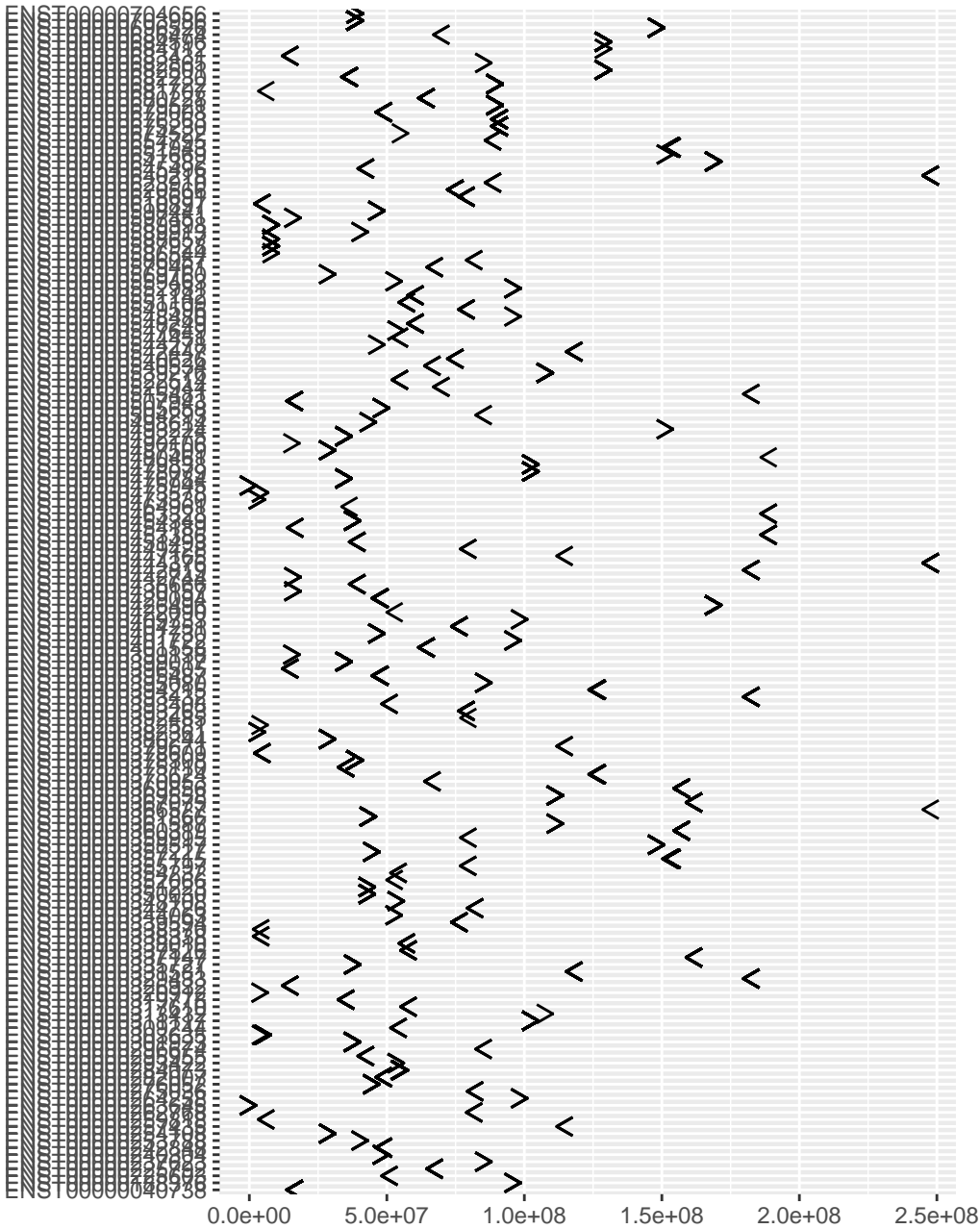


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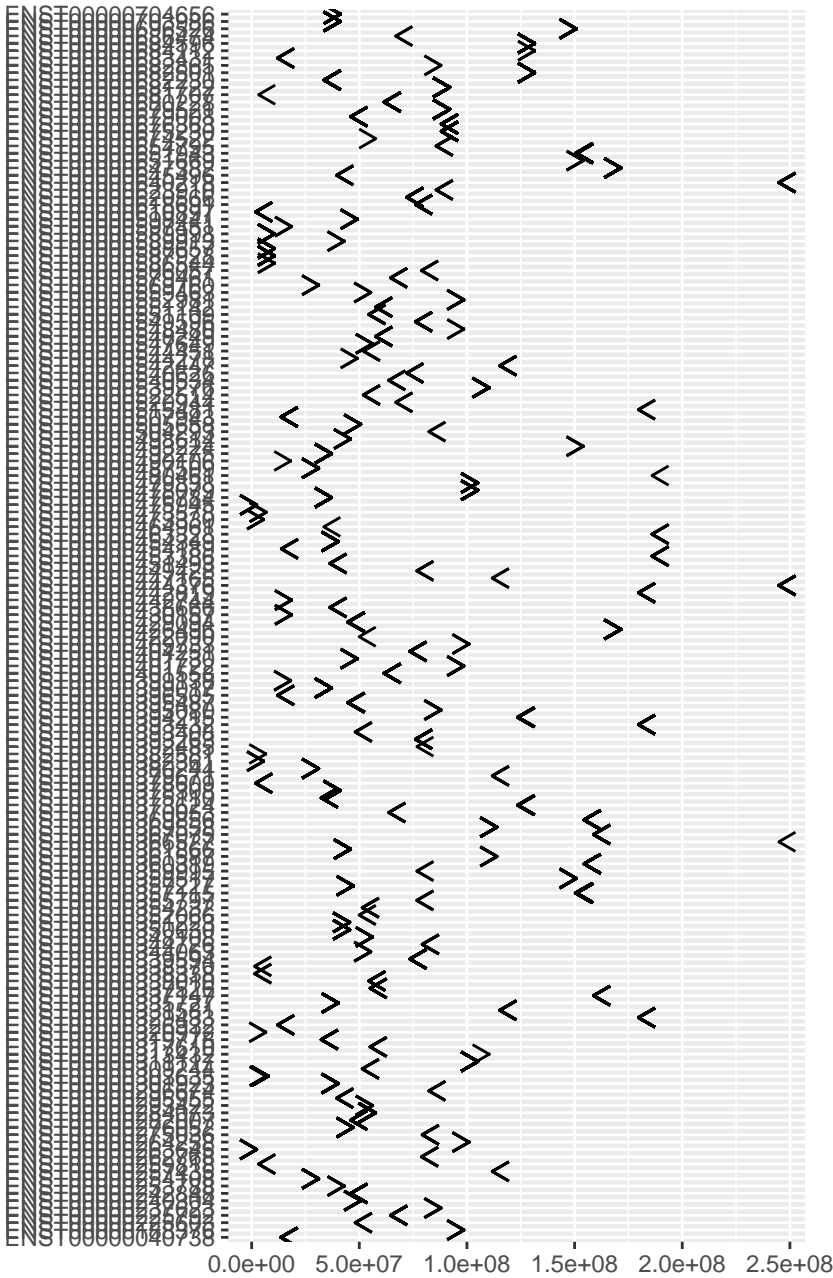
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