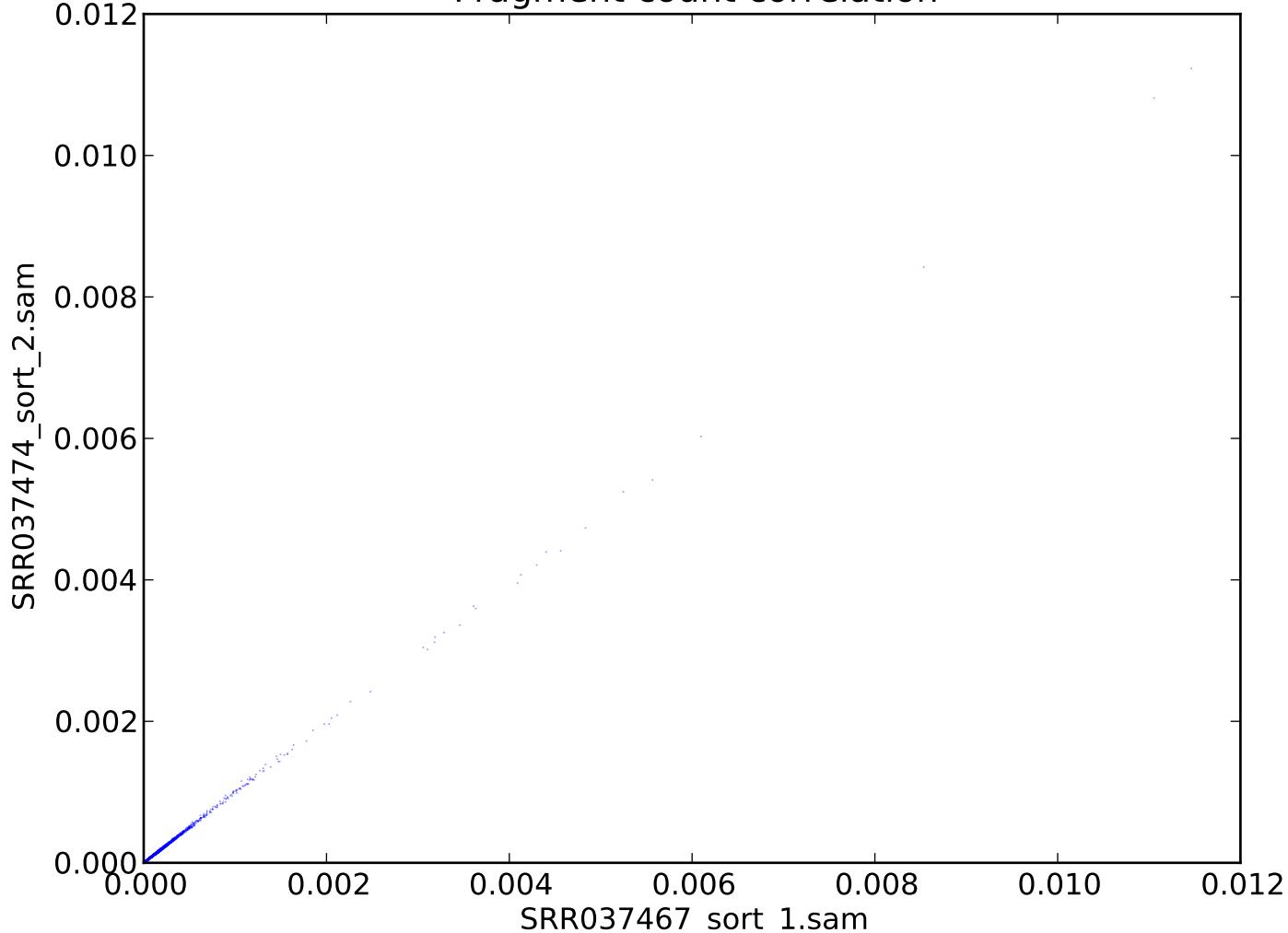
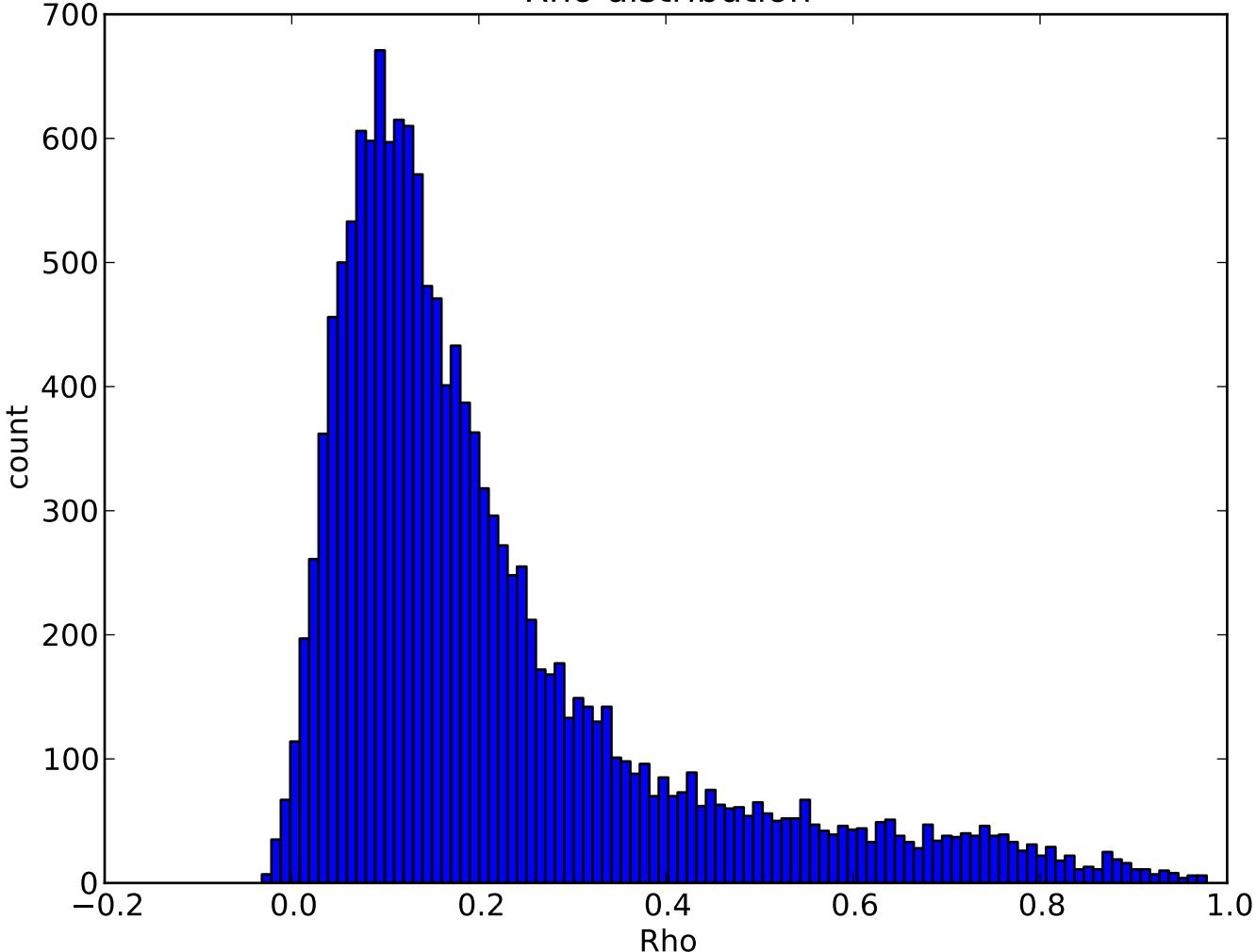


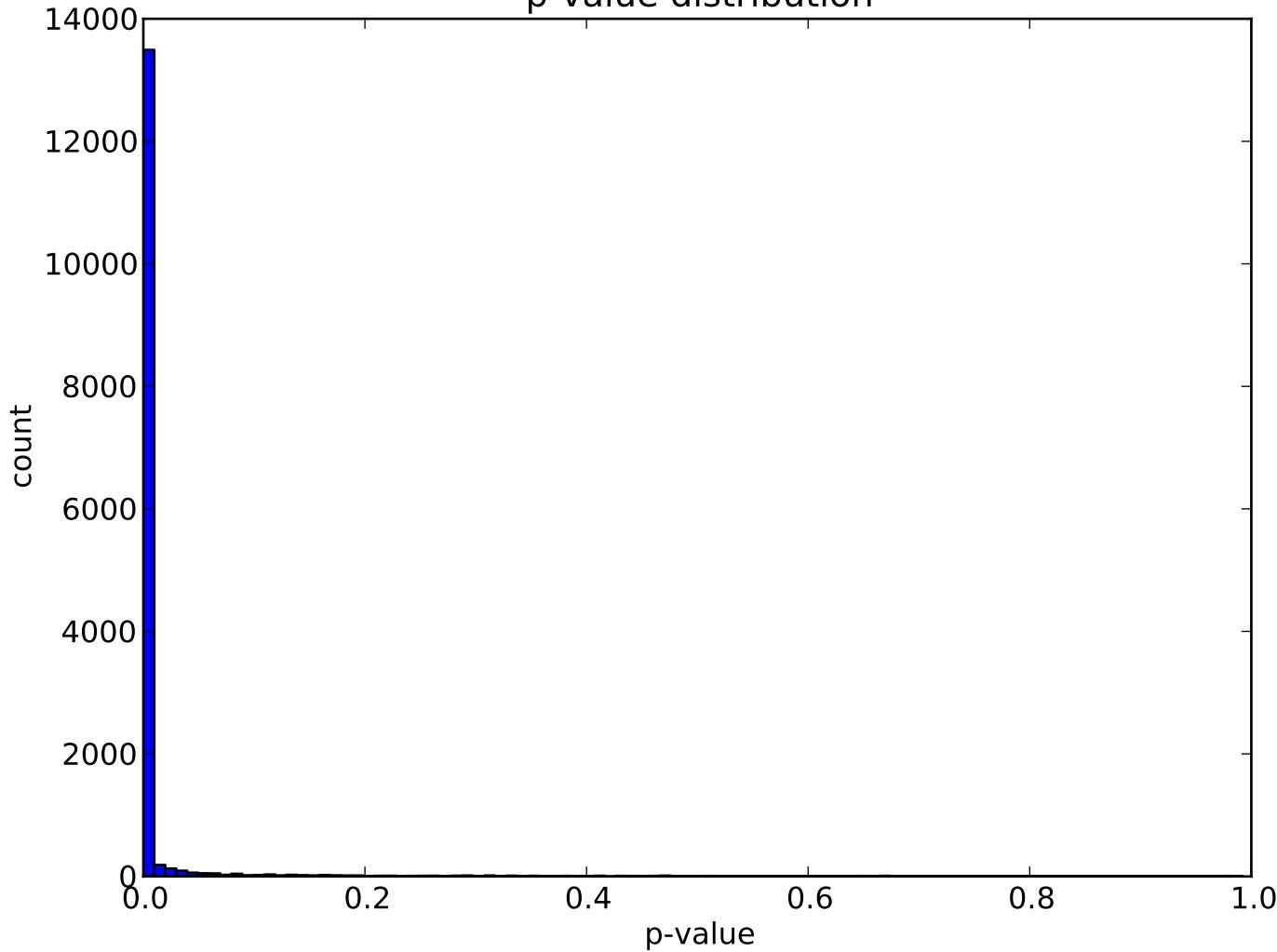
Fragment count correlation



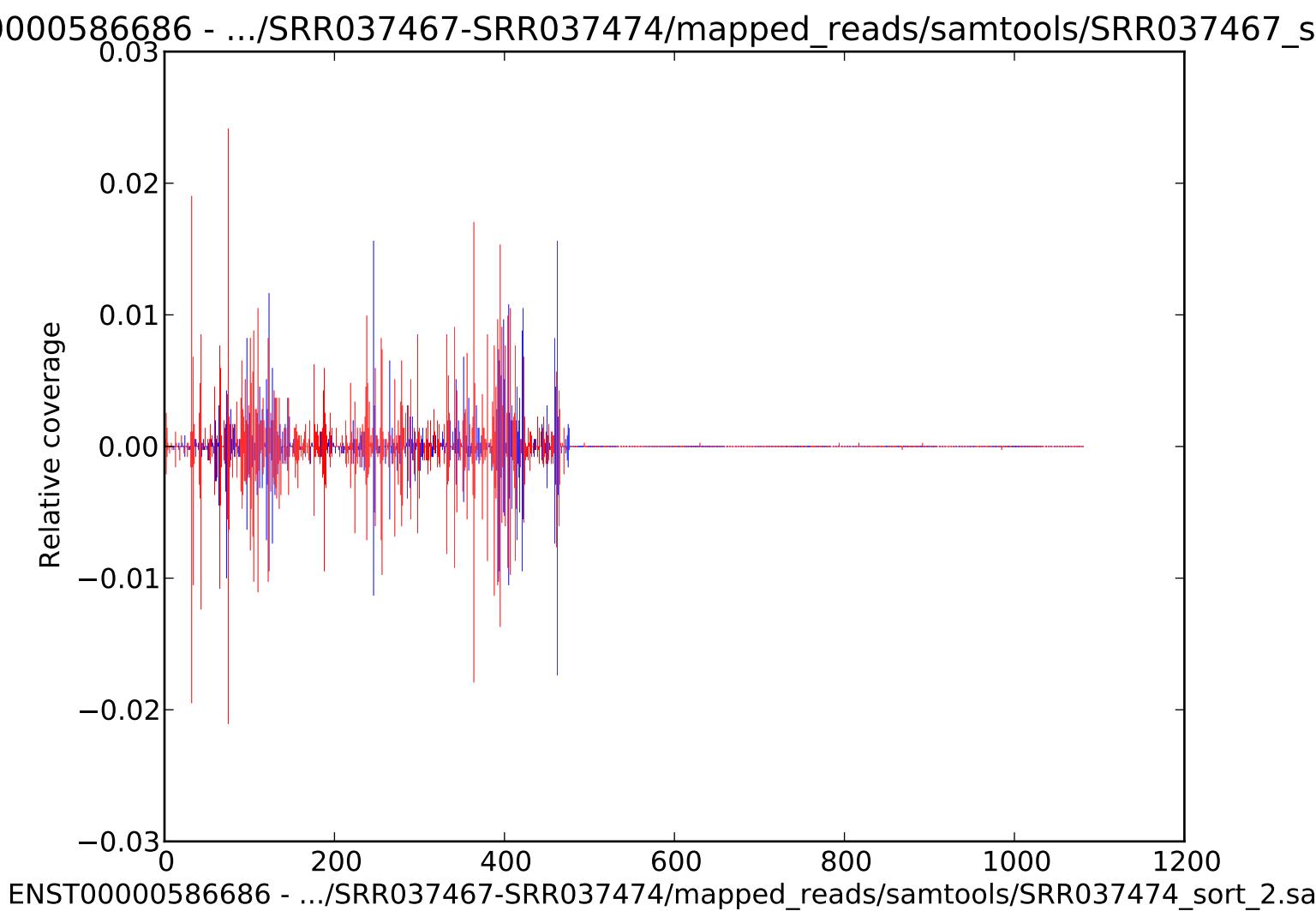
Rho distribution



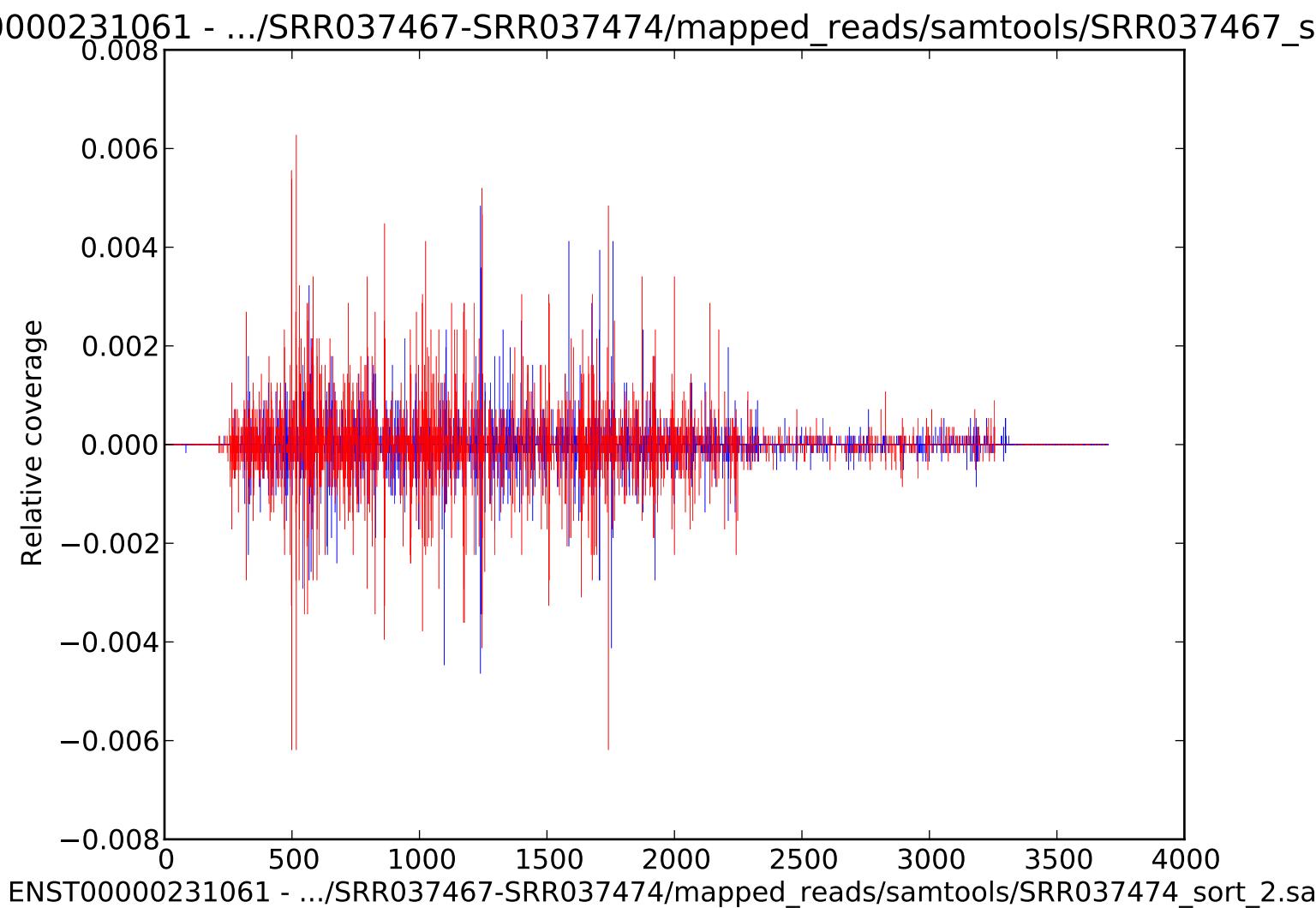
p-value distribution



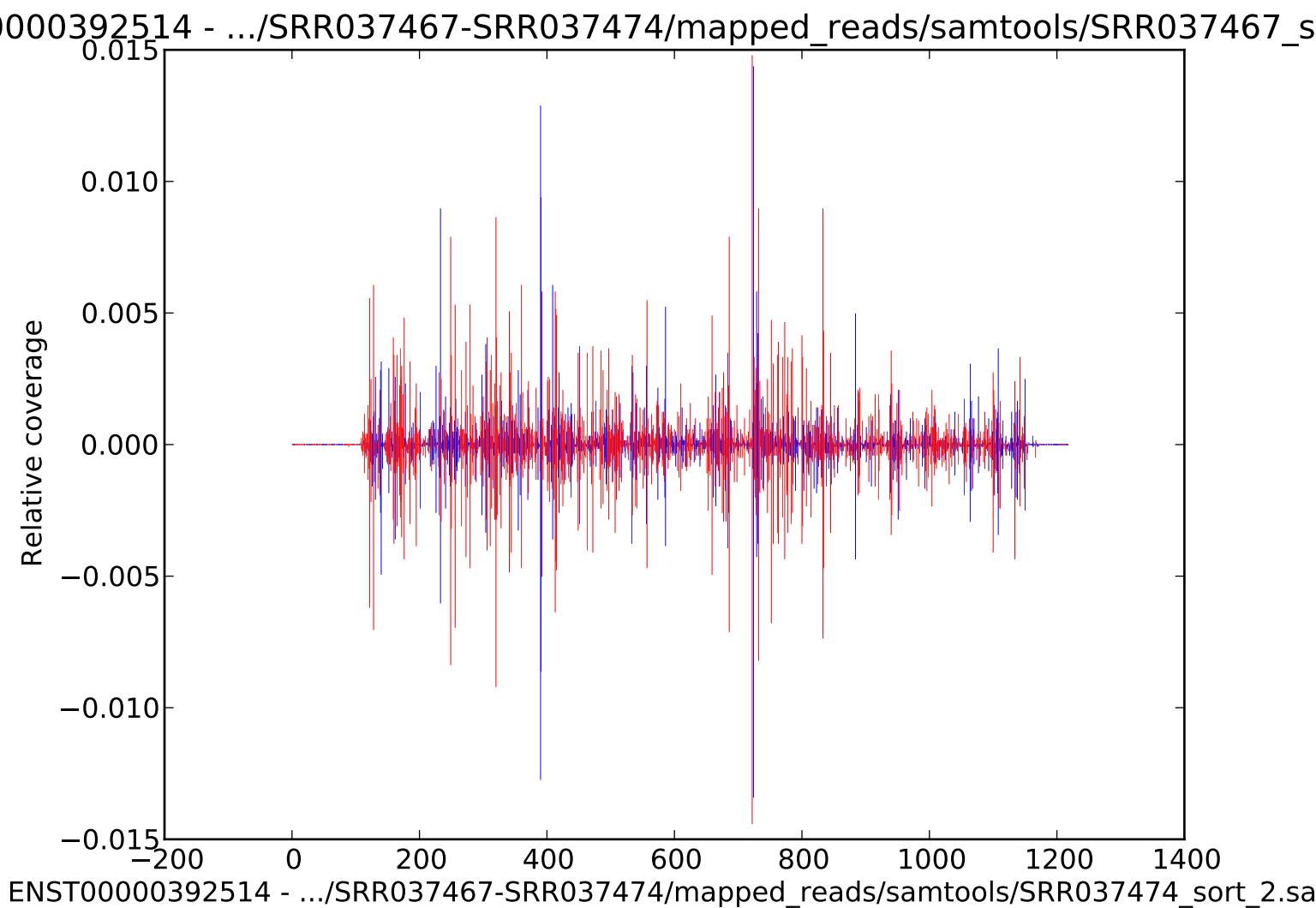
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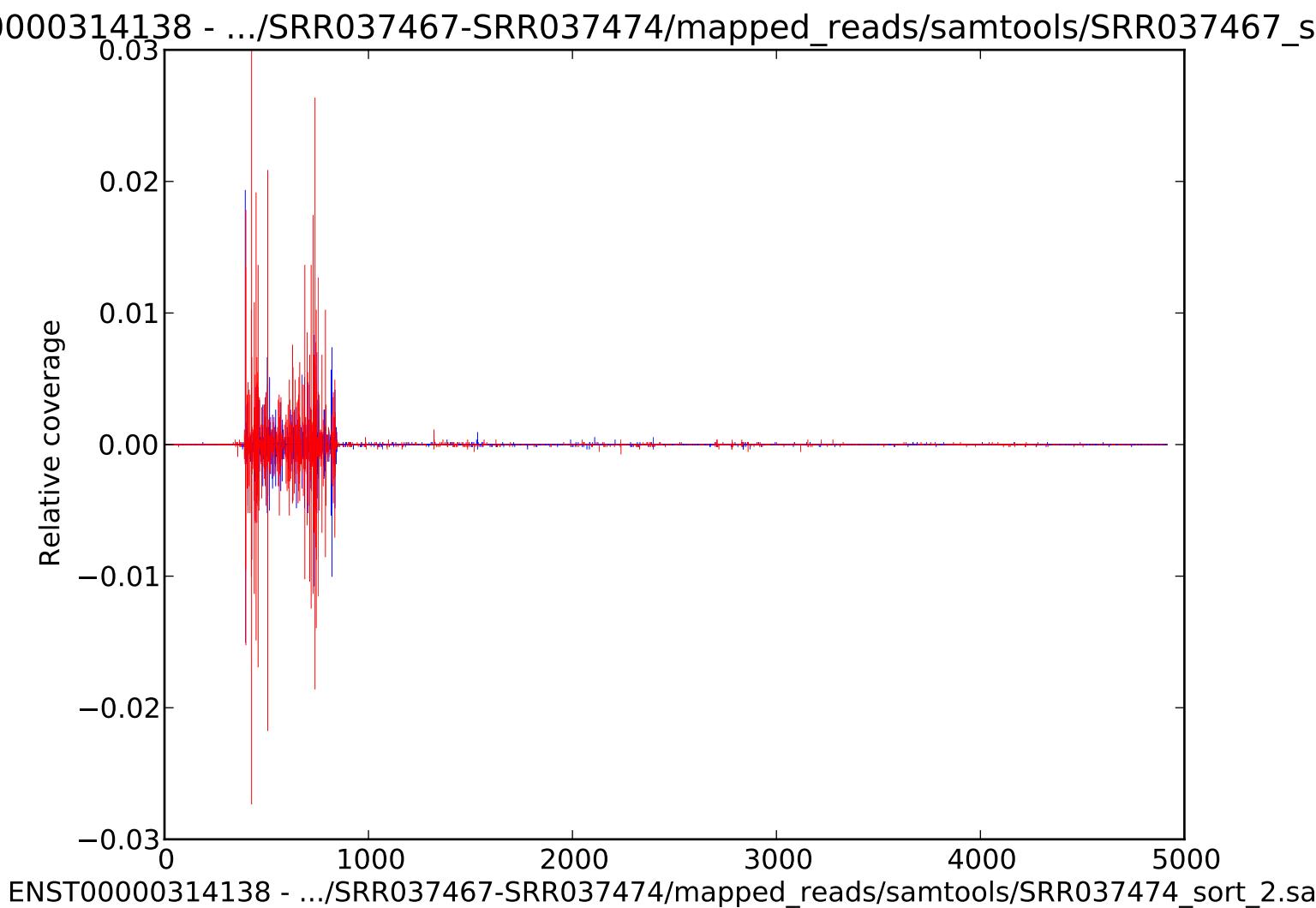
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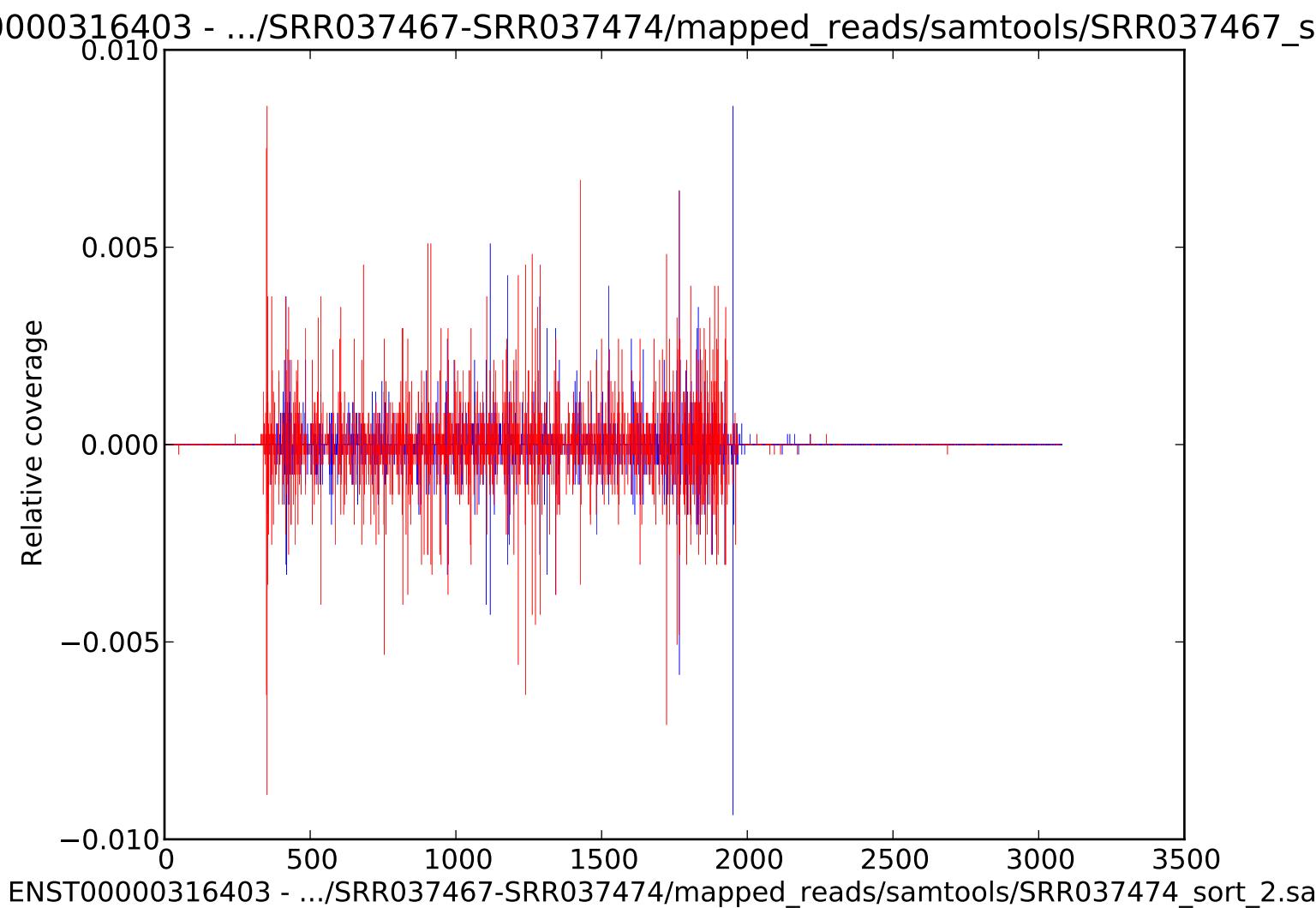
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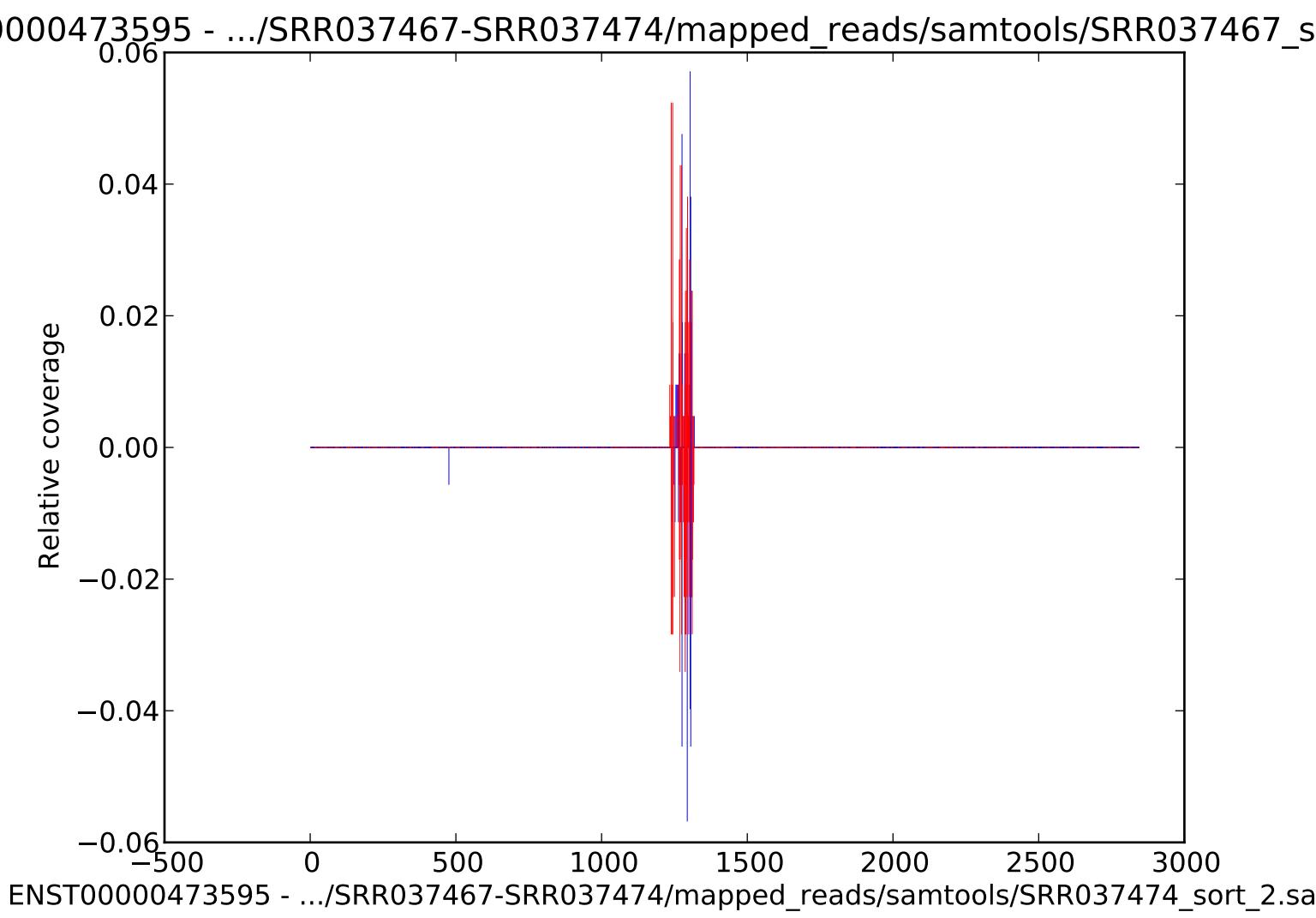
p=0.000000e+00, Rho=7.042373e-01



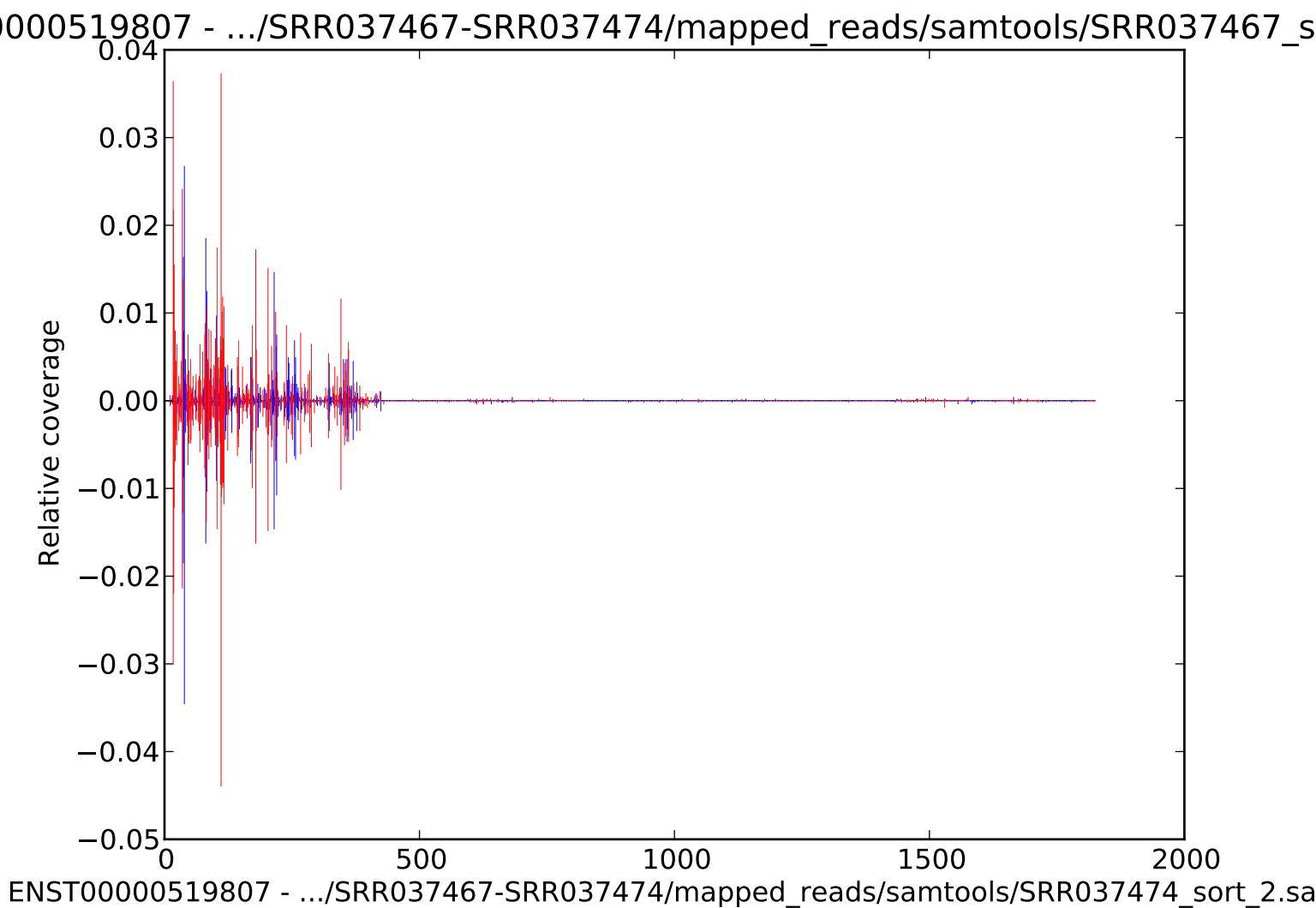
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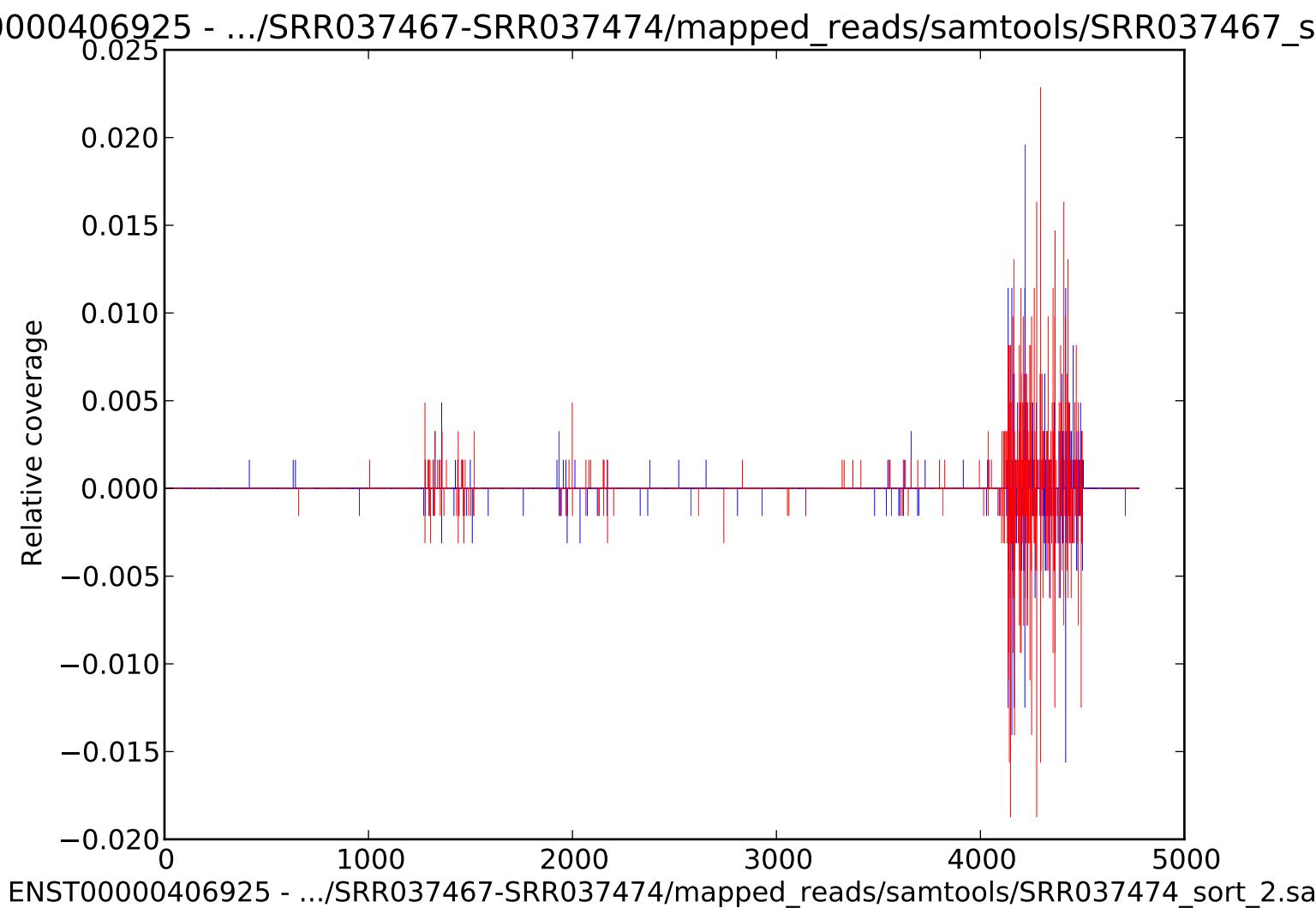
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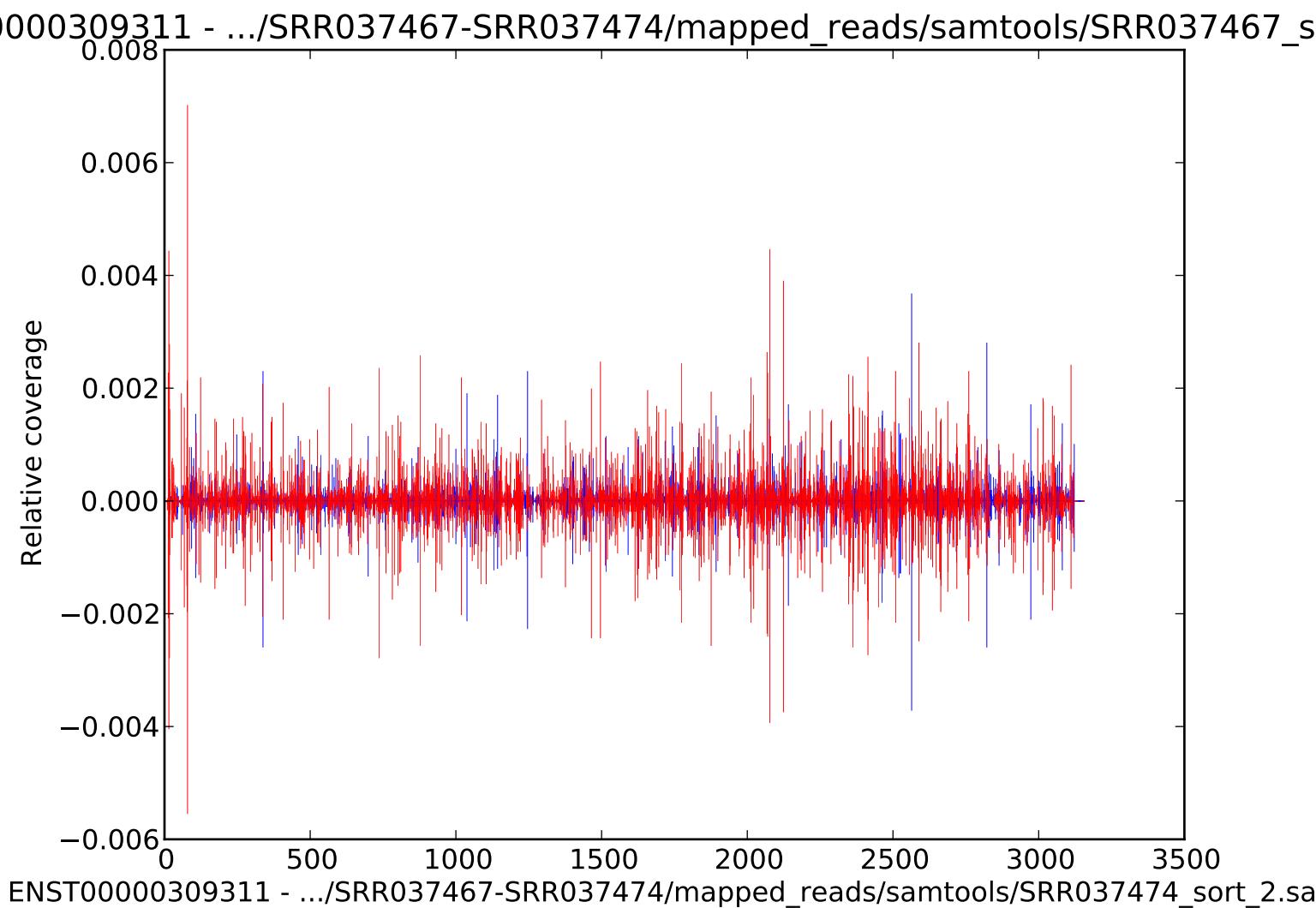
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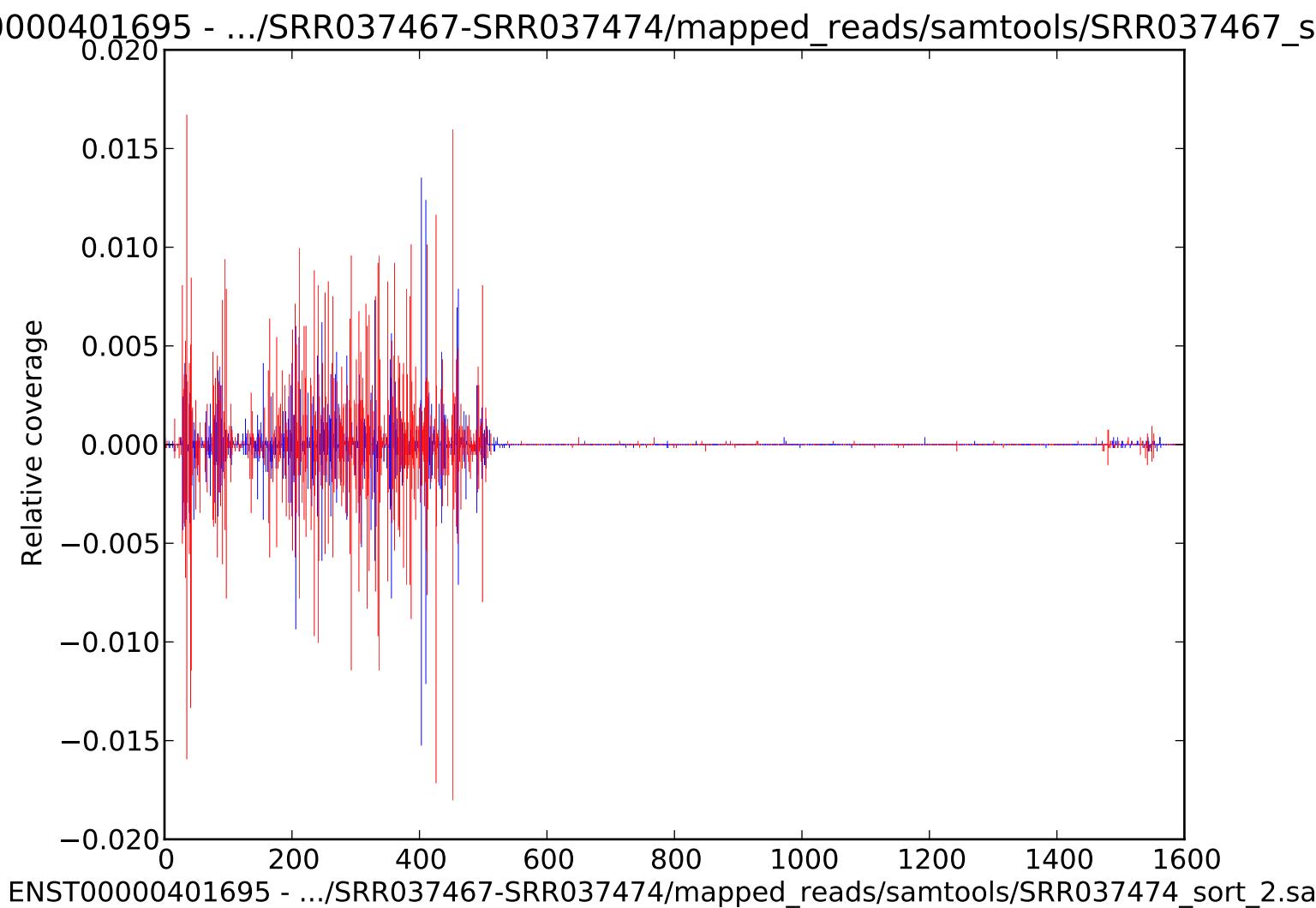
p=0.000000e+00, Rho=5.101216e-01



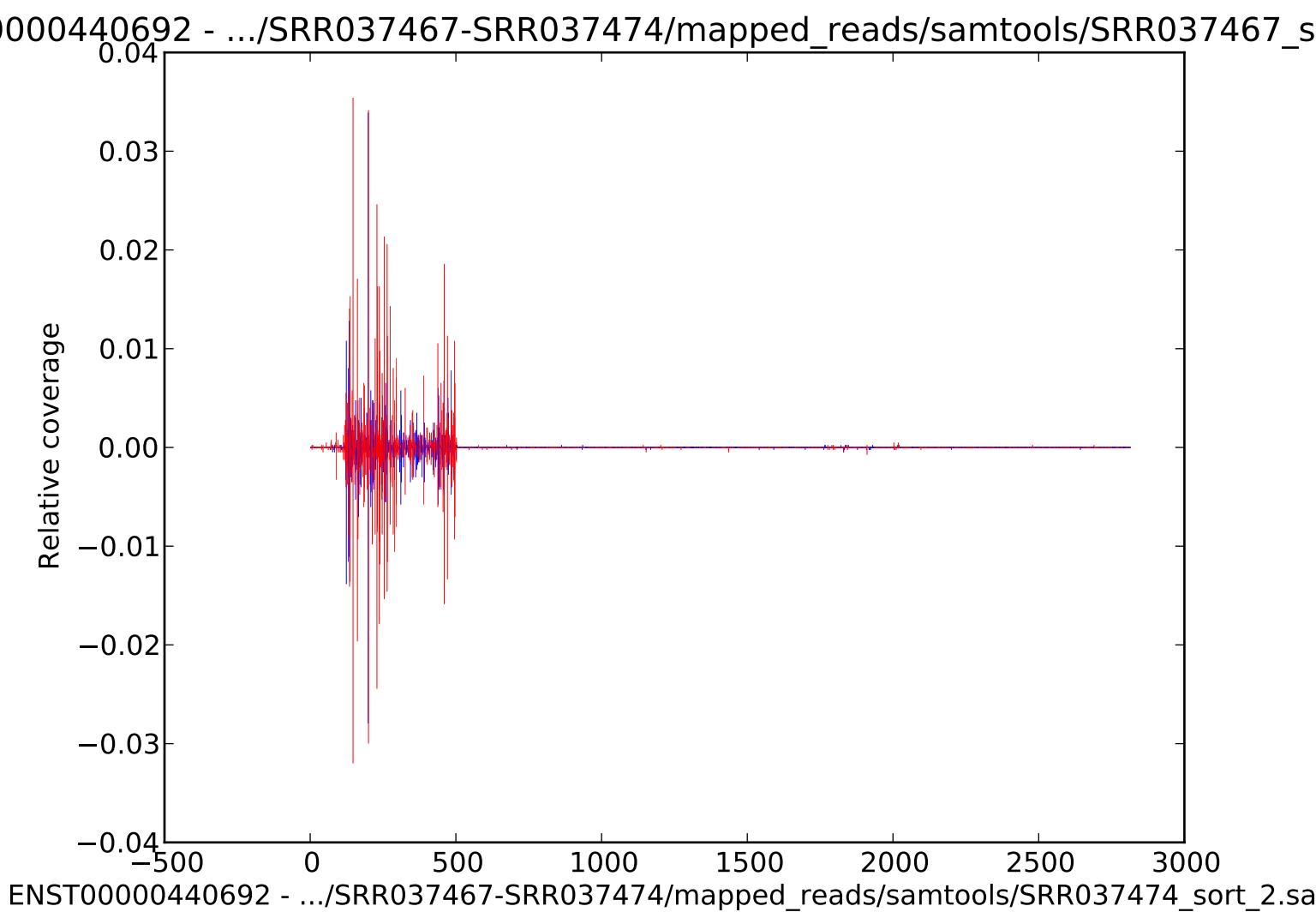
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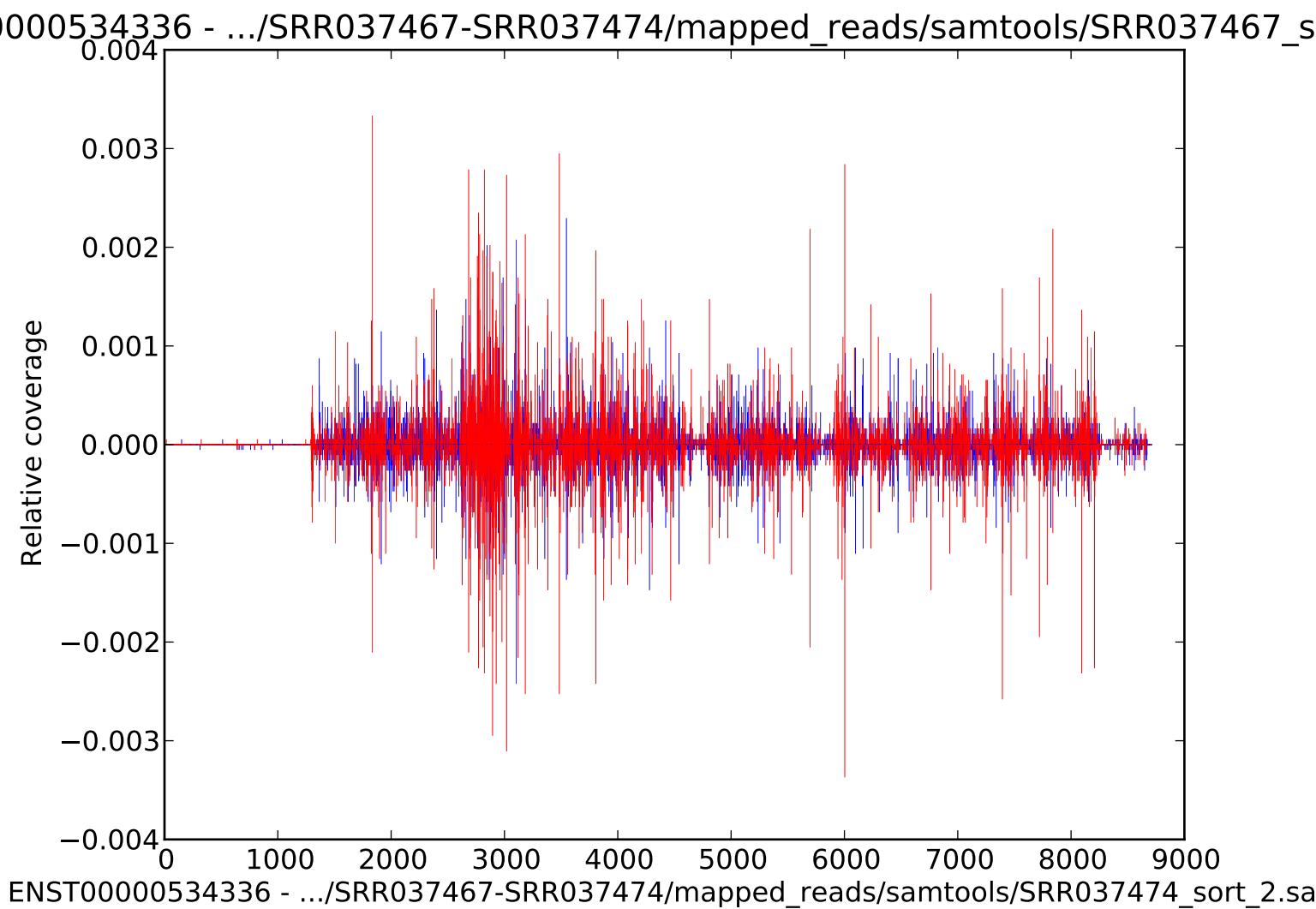
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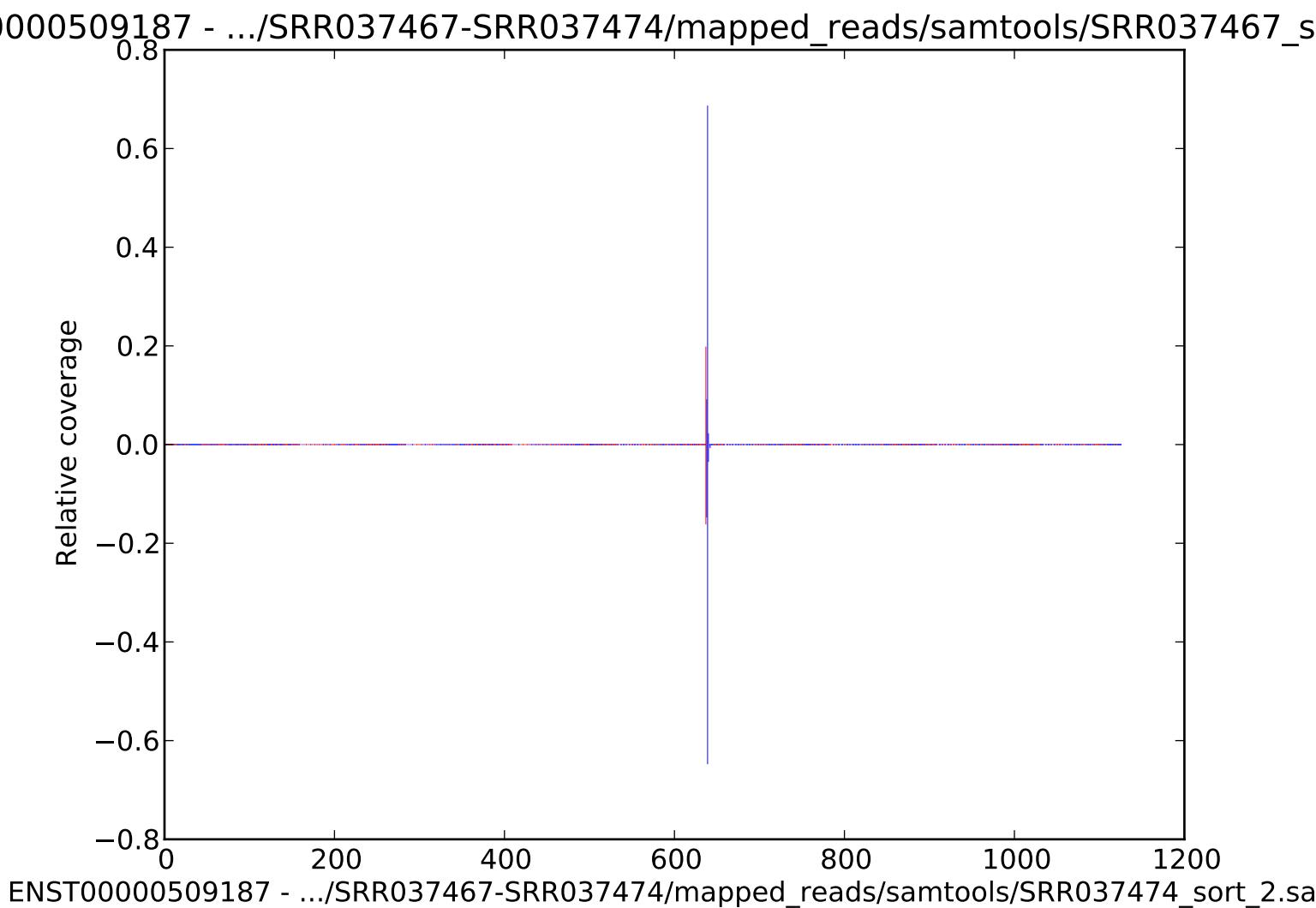
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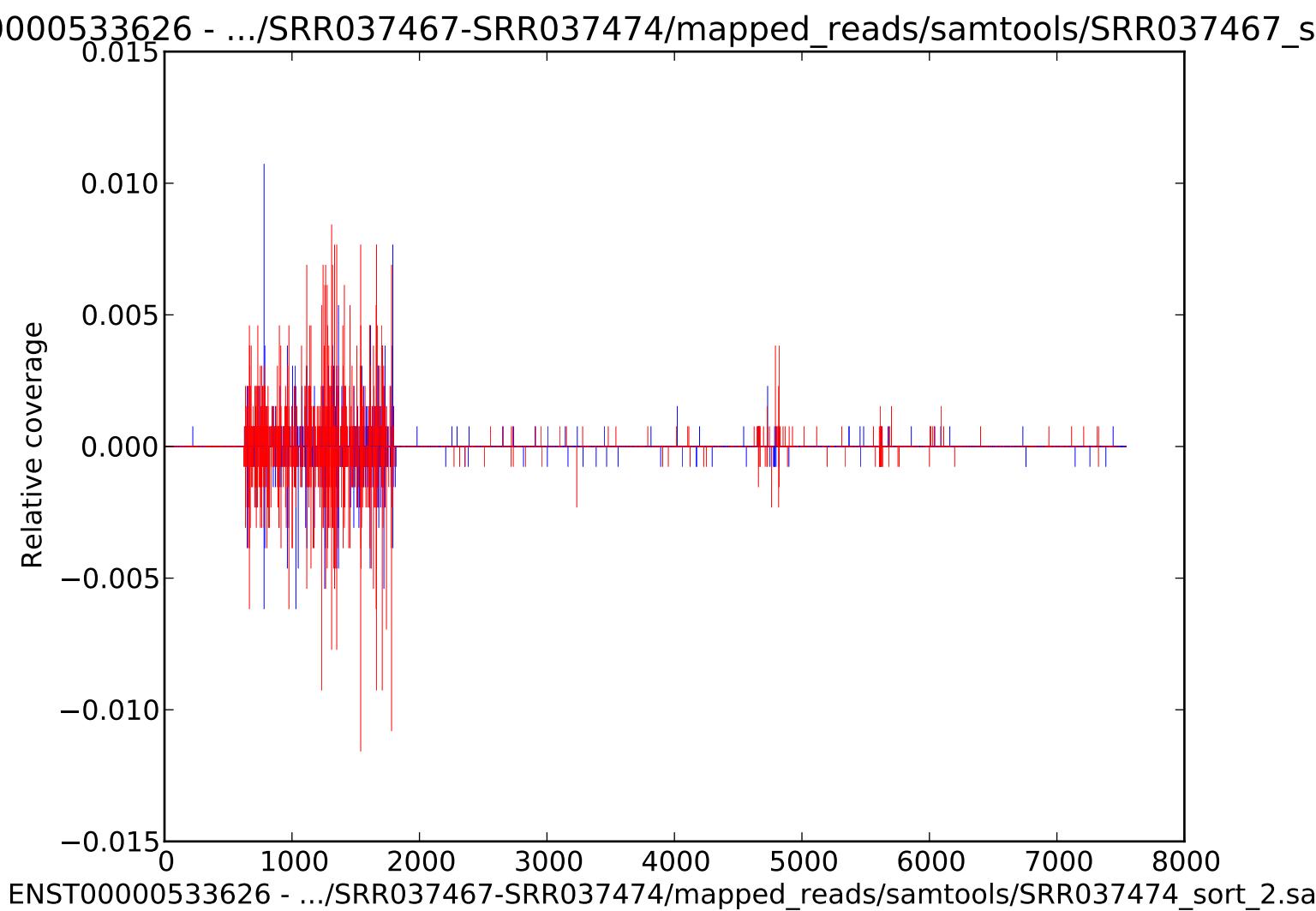
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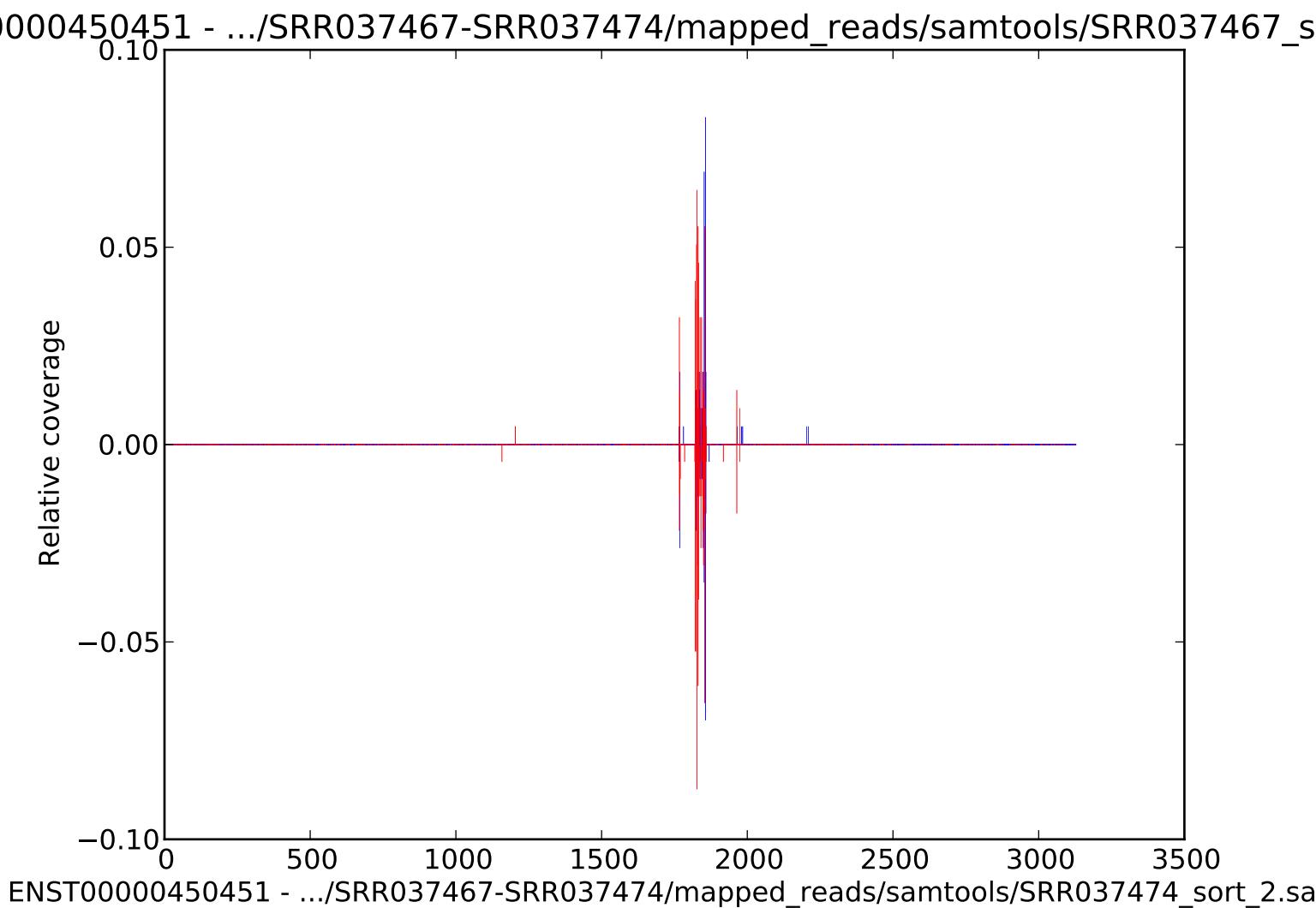
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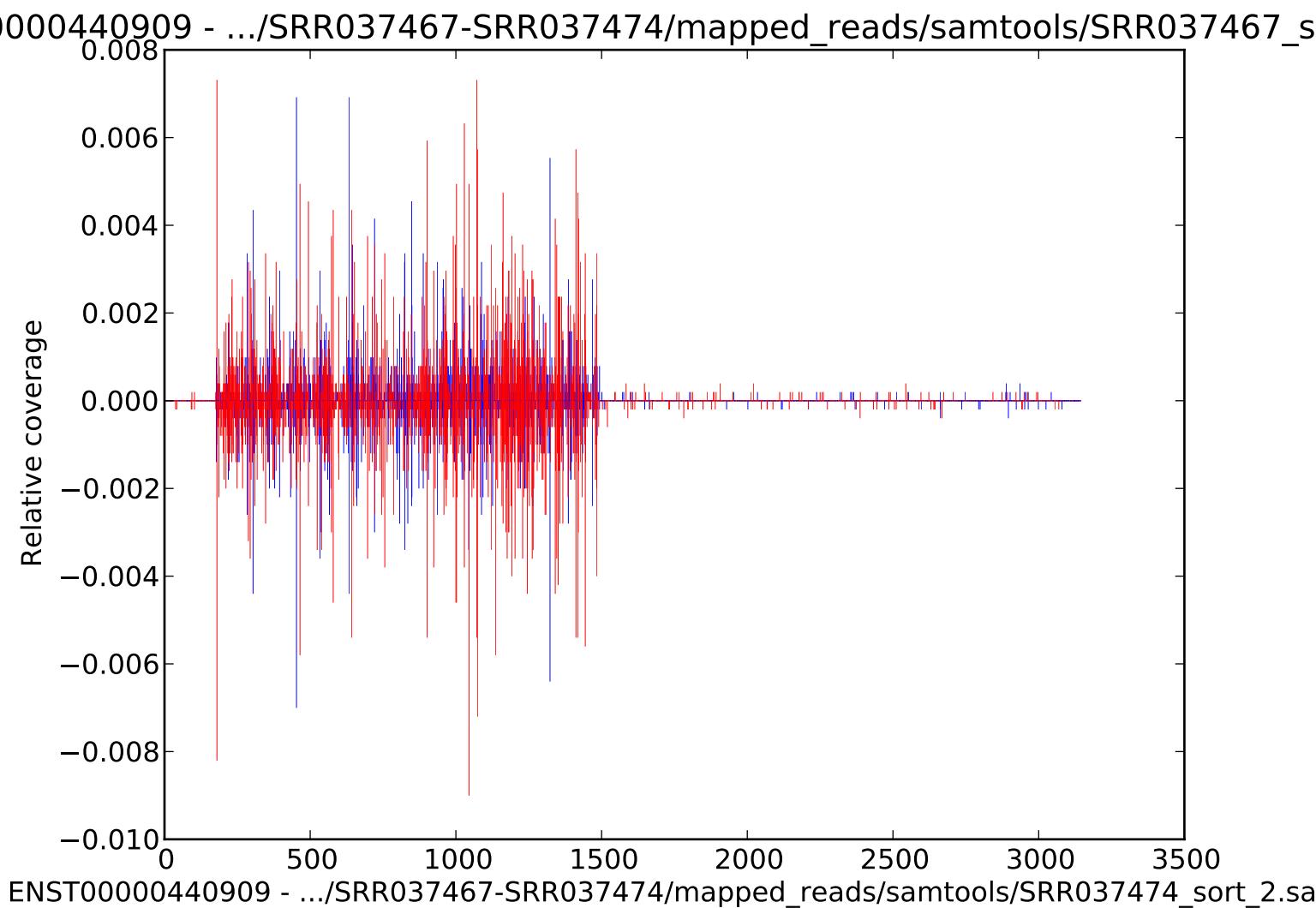
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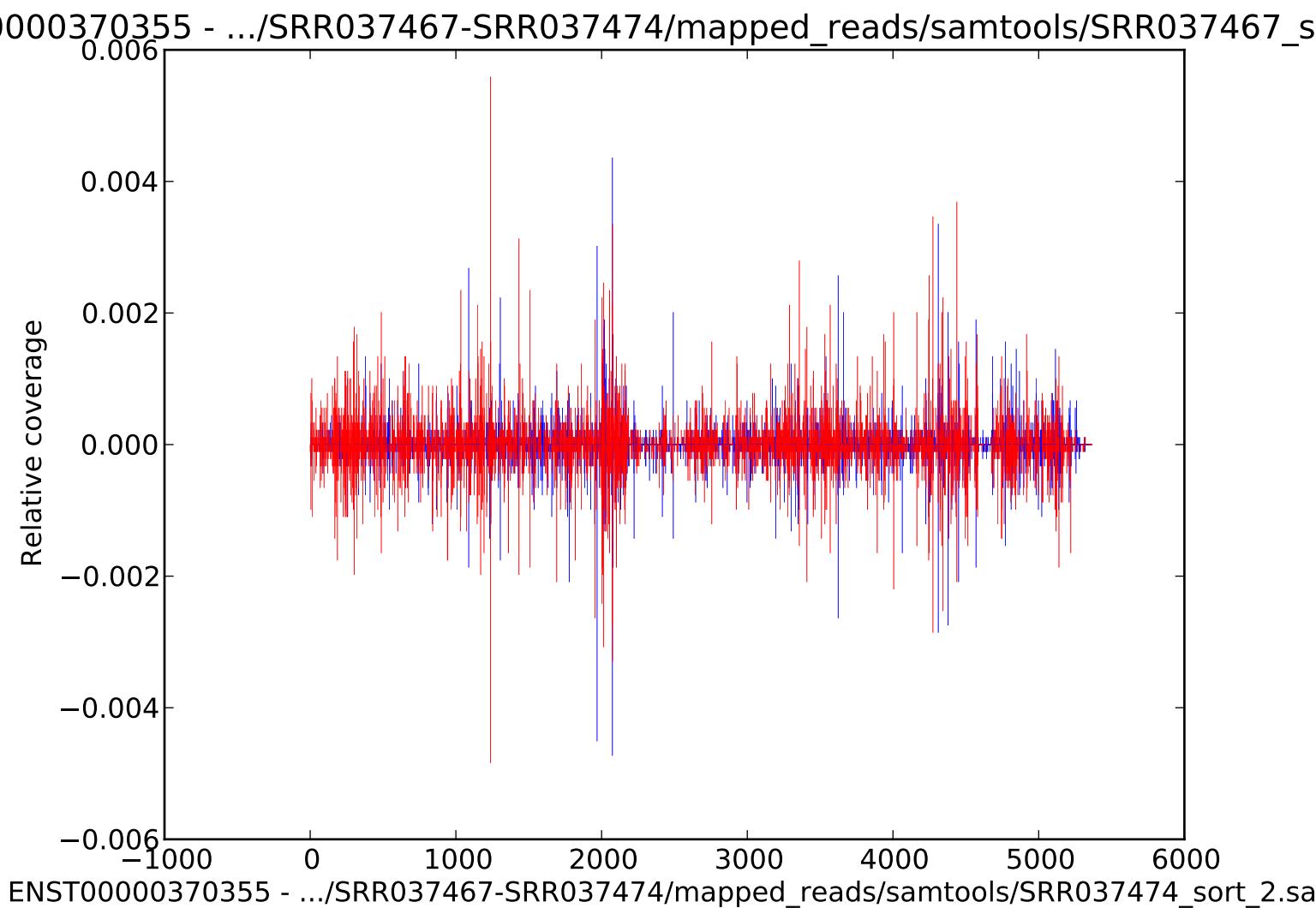
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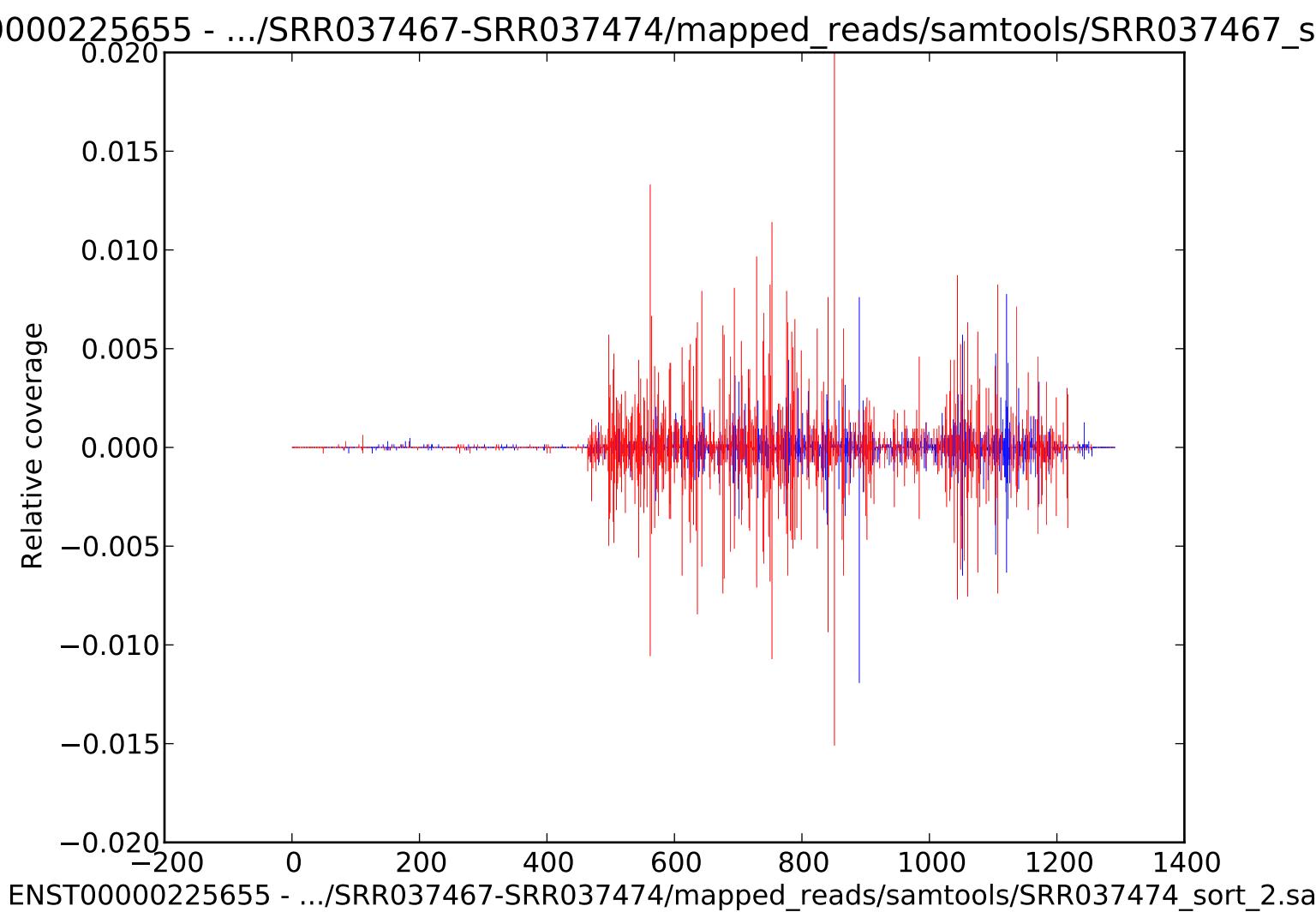
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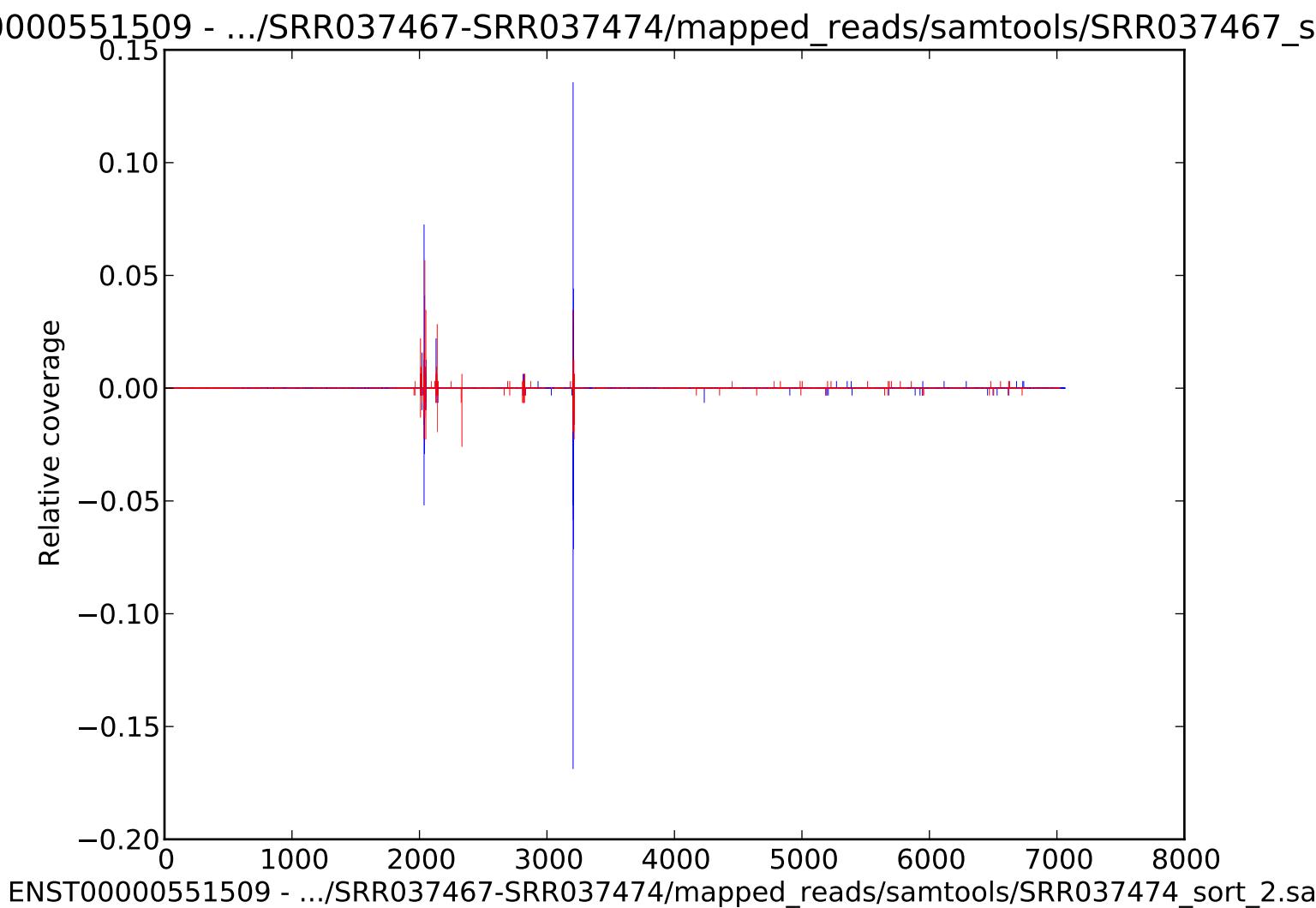
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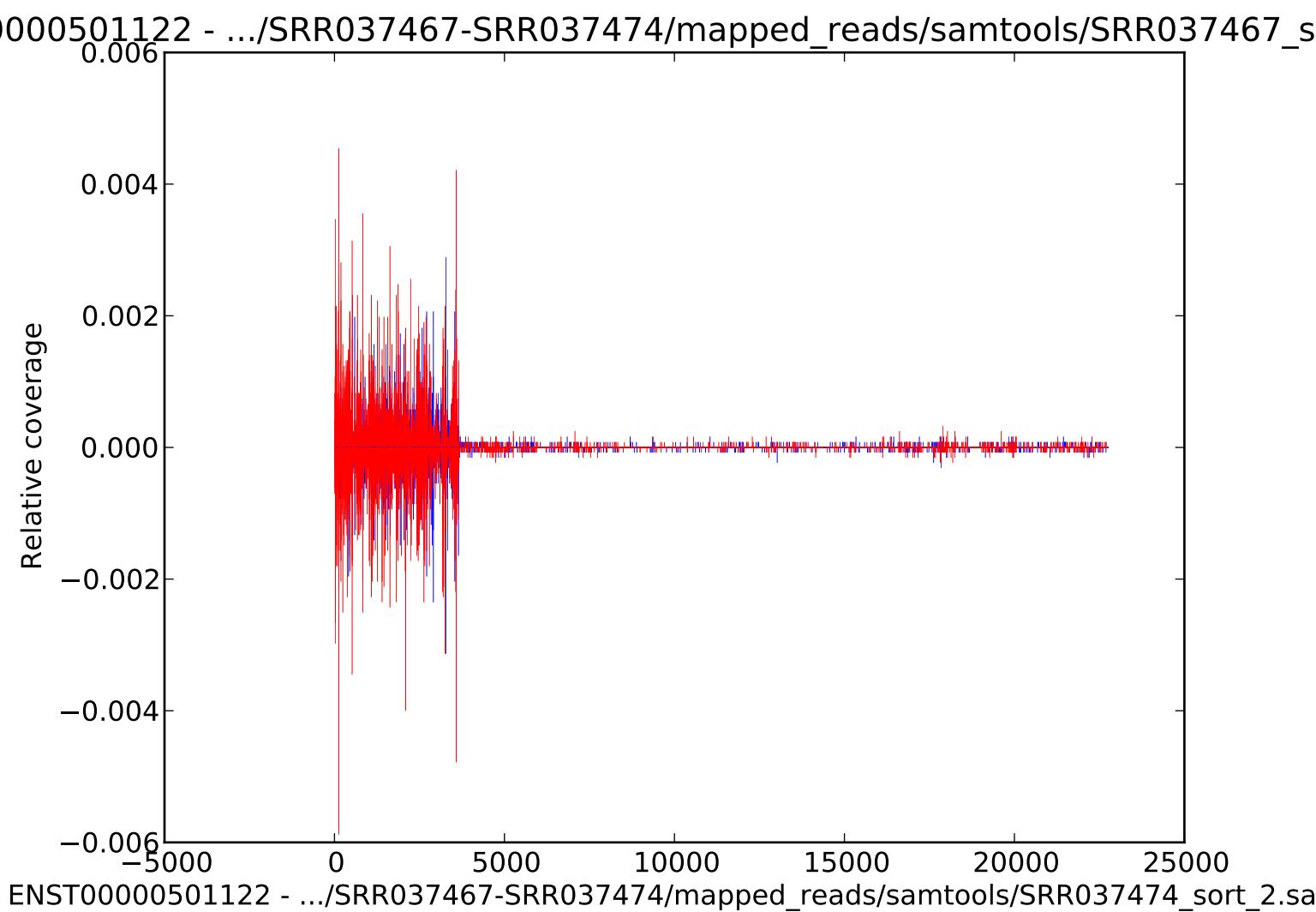
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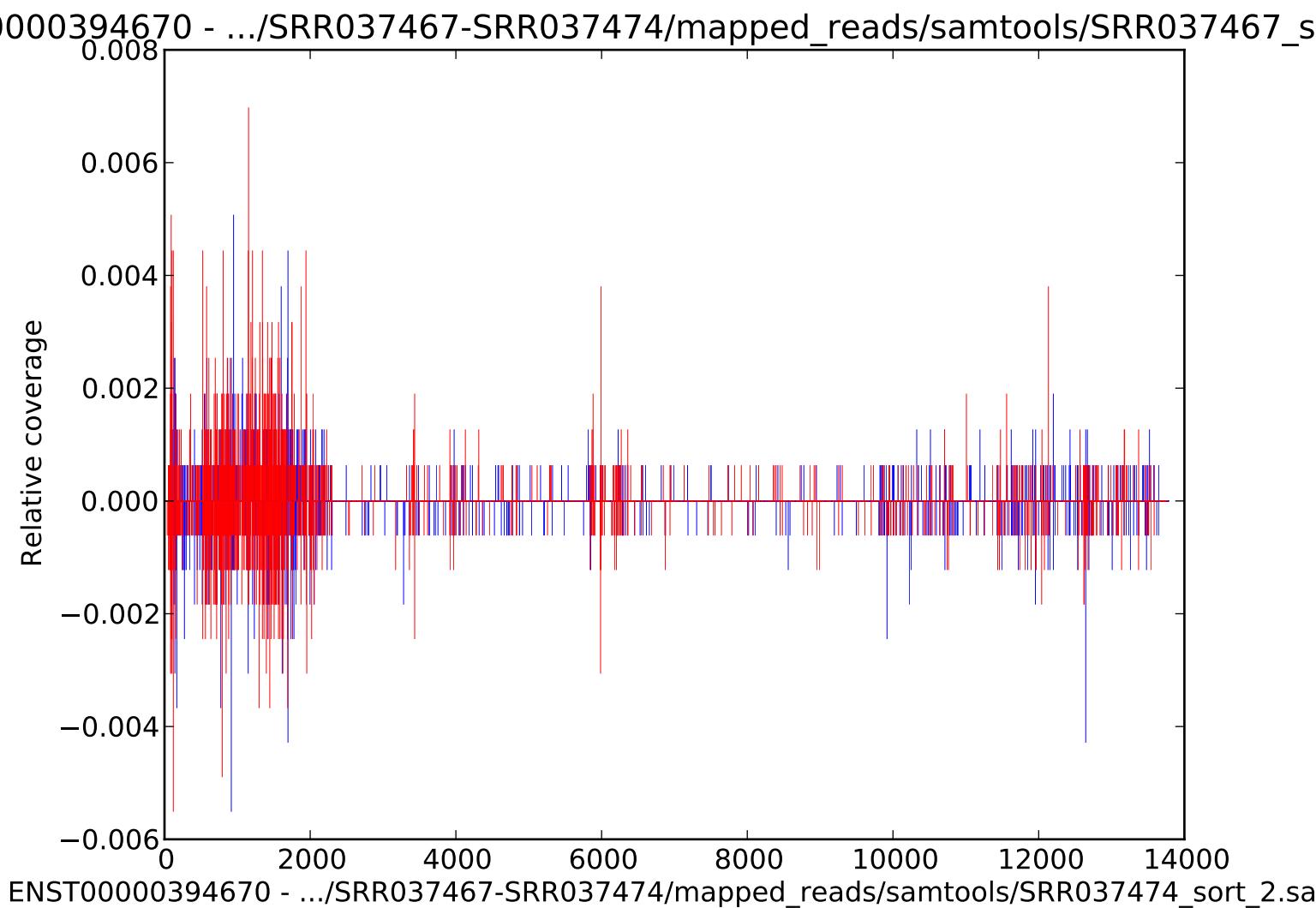
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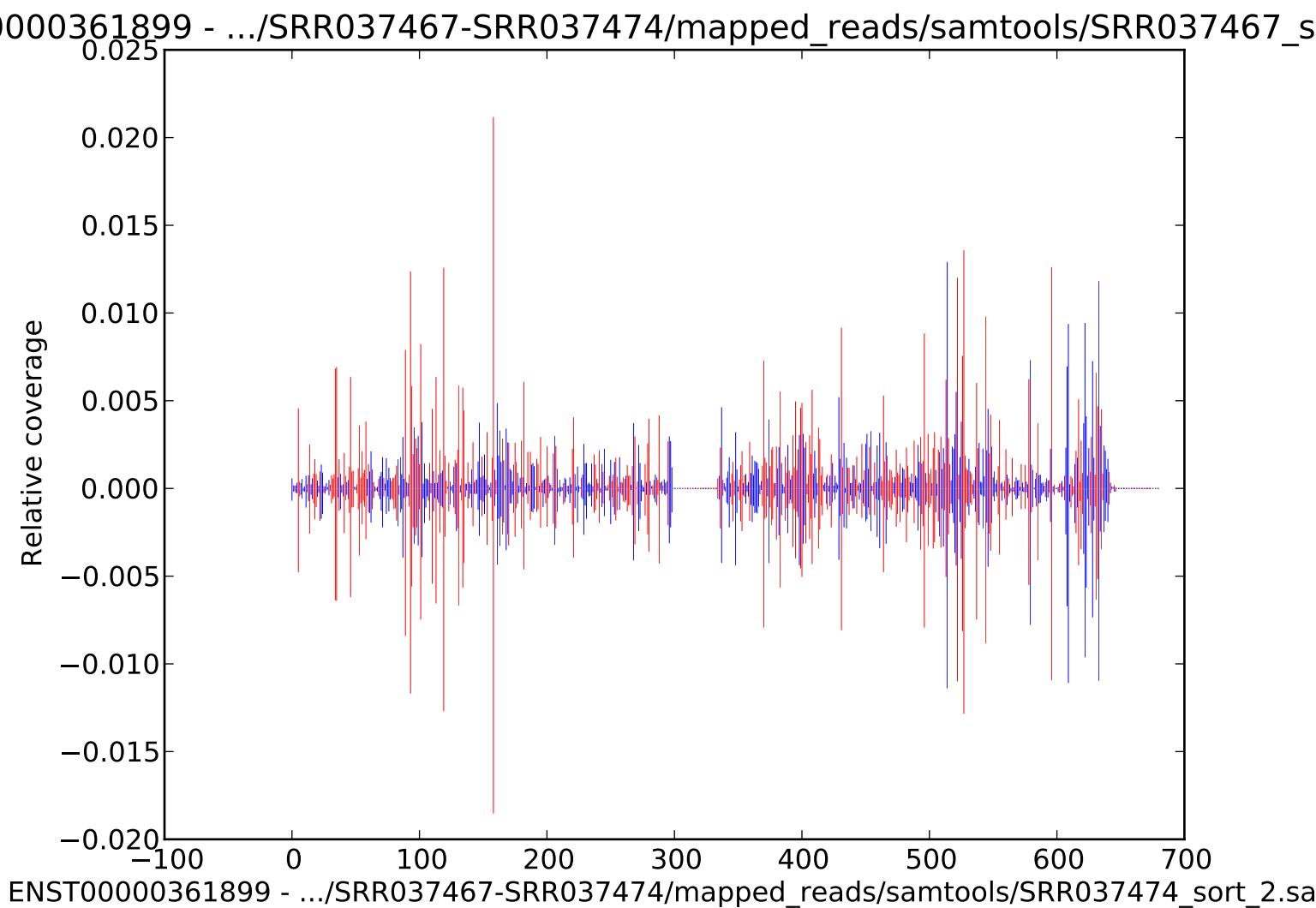
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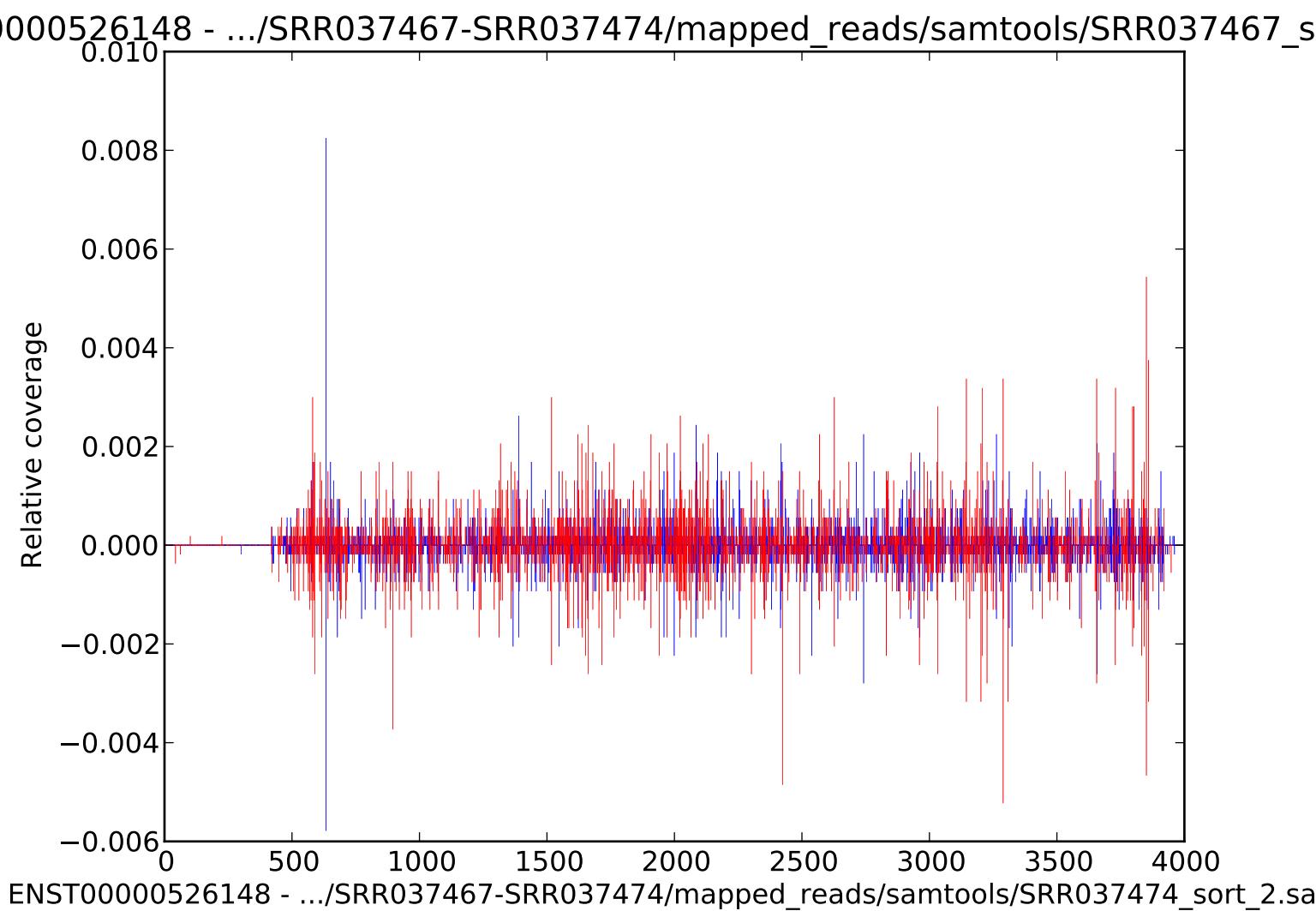
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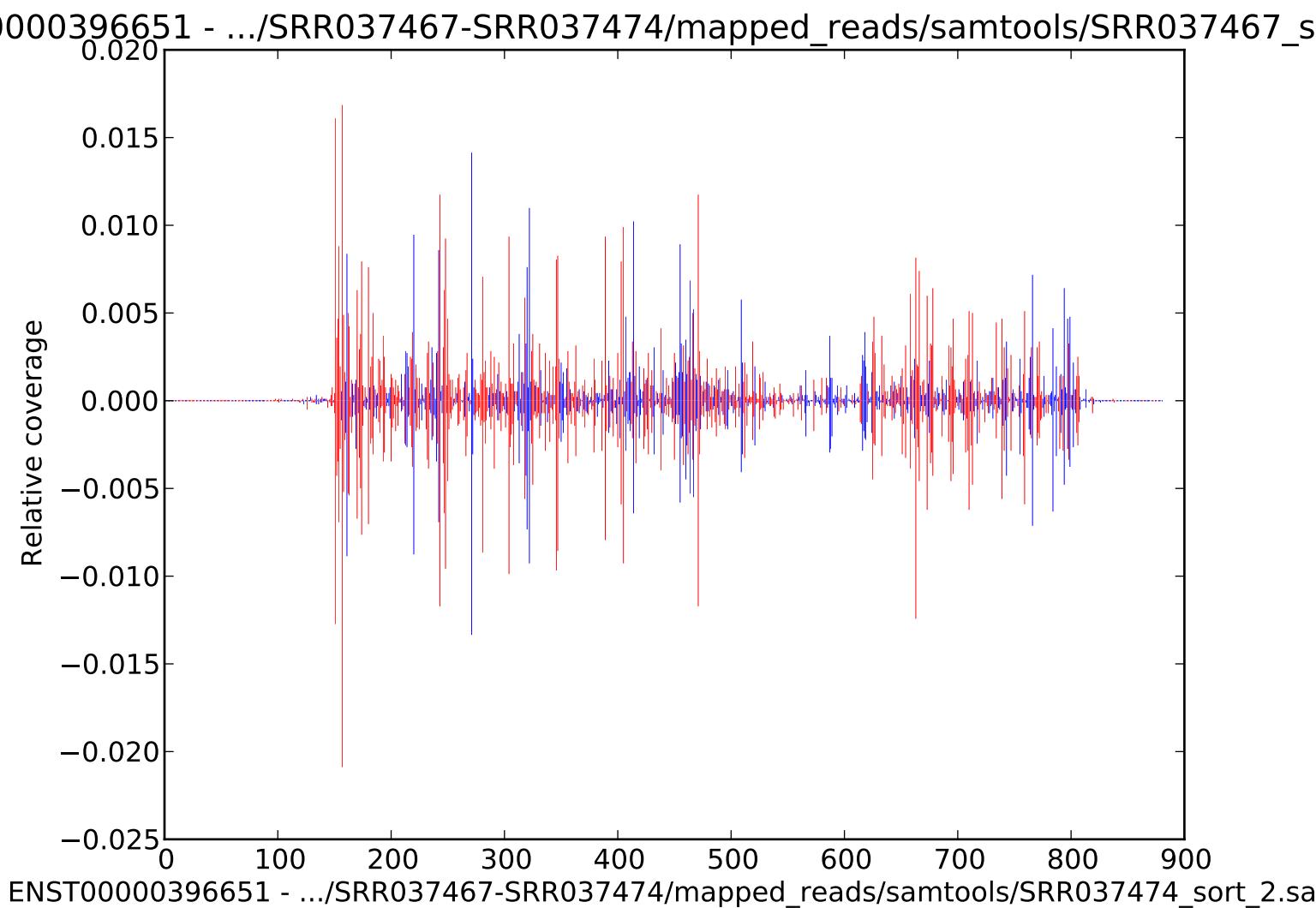
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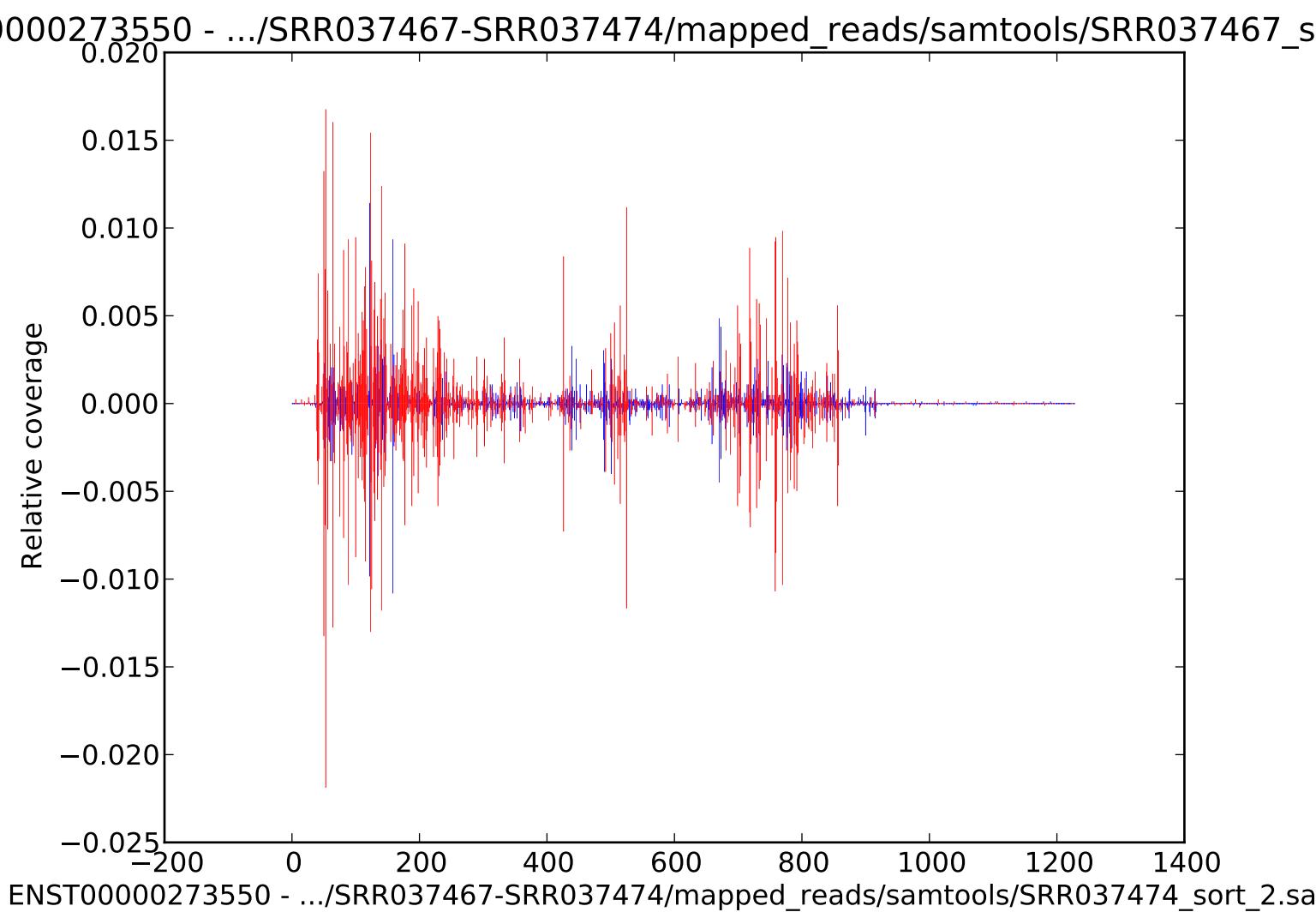
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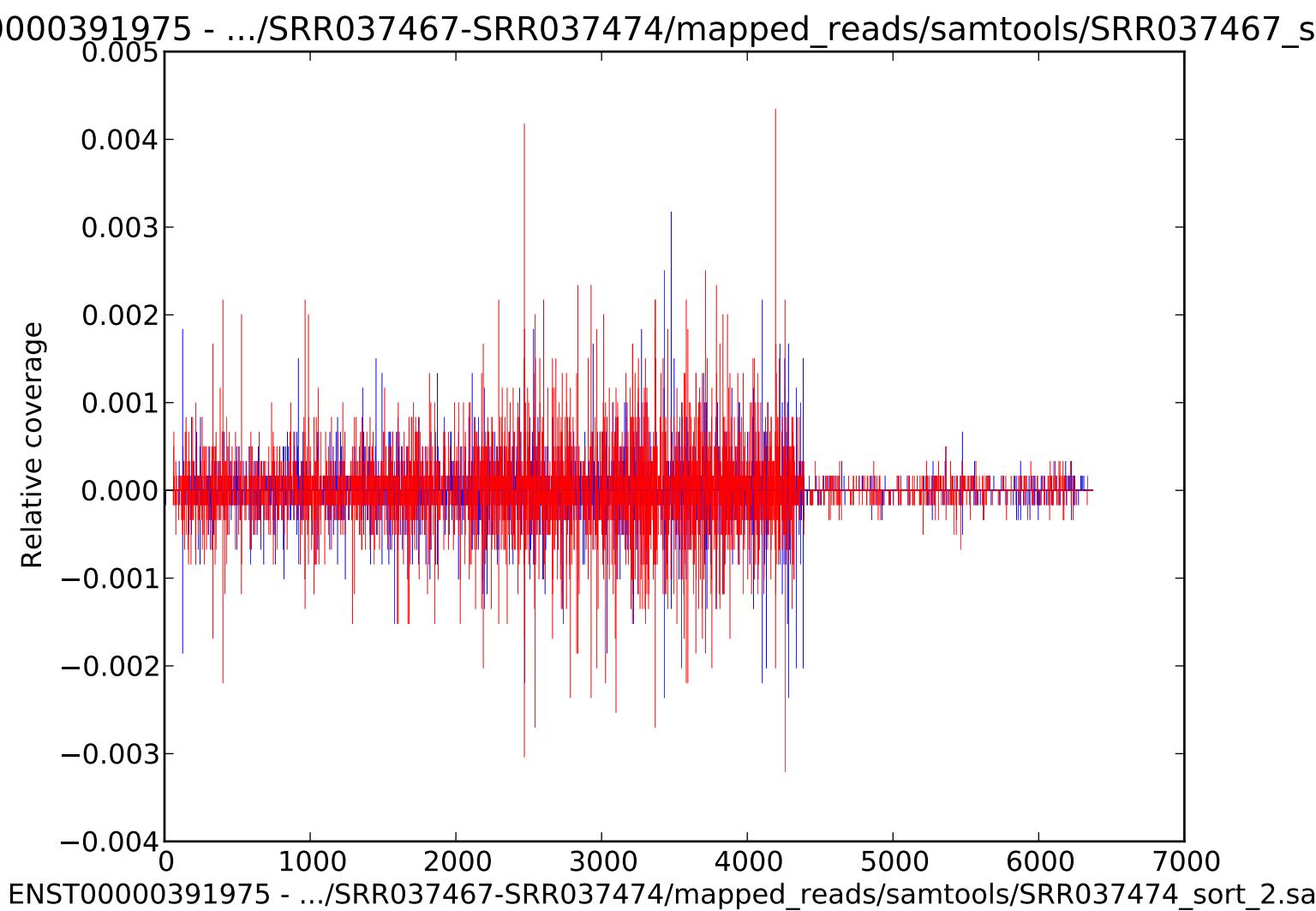
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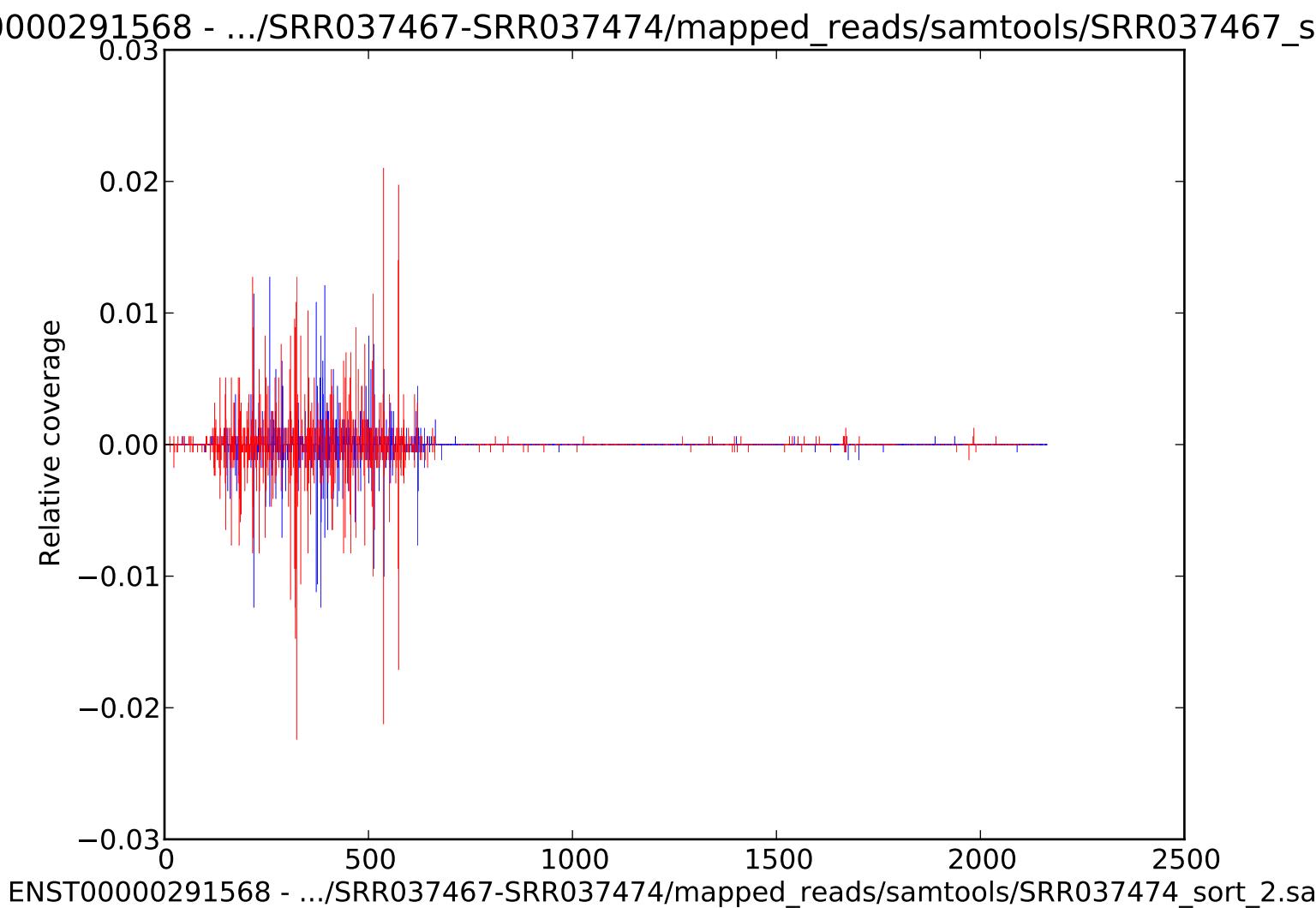
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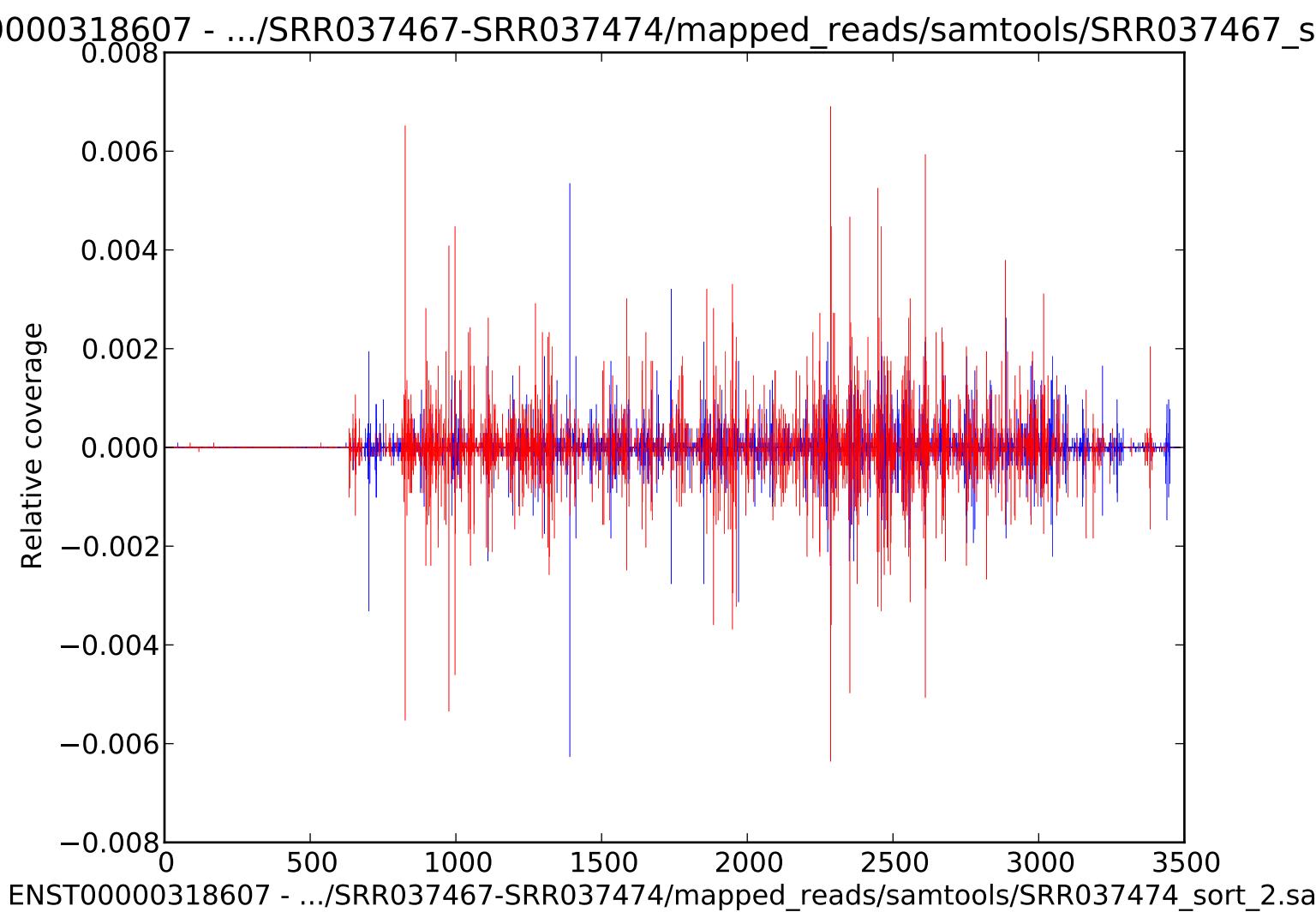
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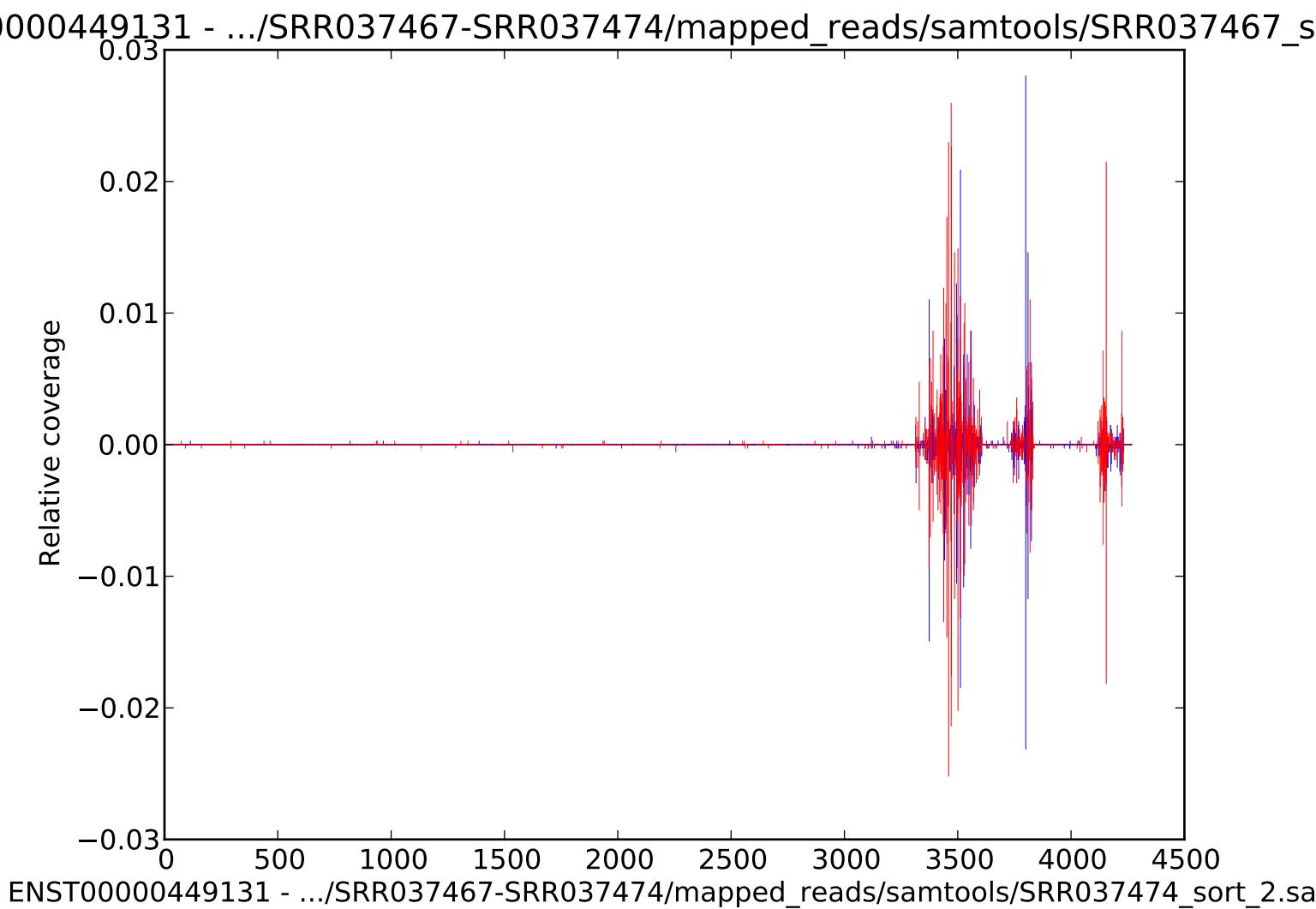
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p=0.000000e+00, Rho=7.887483e-01



p=0.000000e+00, Rho=7.791994e-01



p=0.000000e+00, Rho=7.874633e-01

