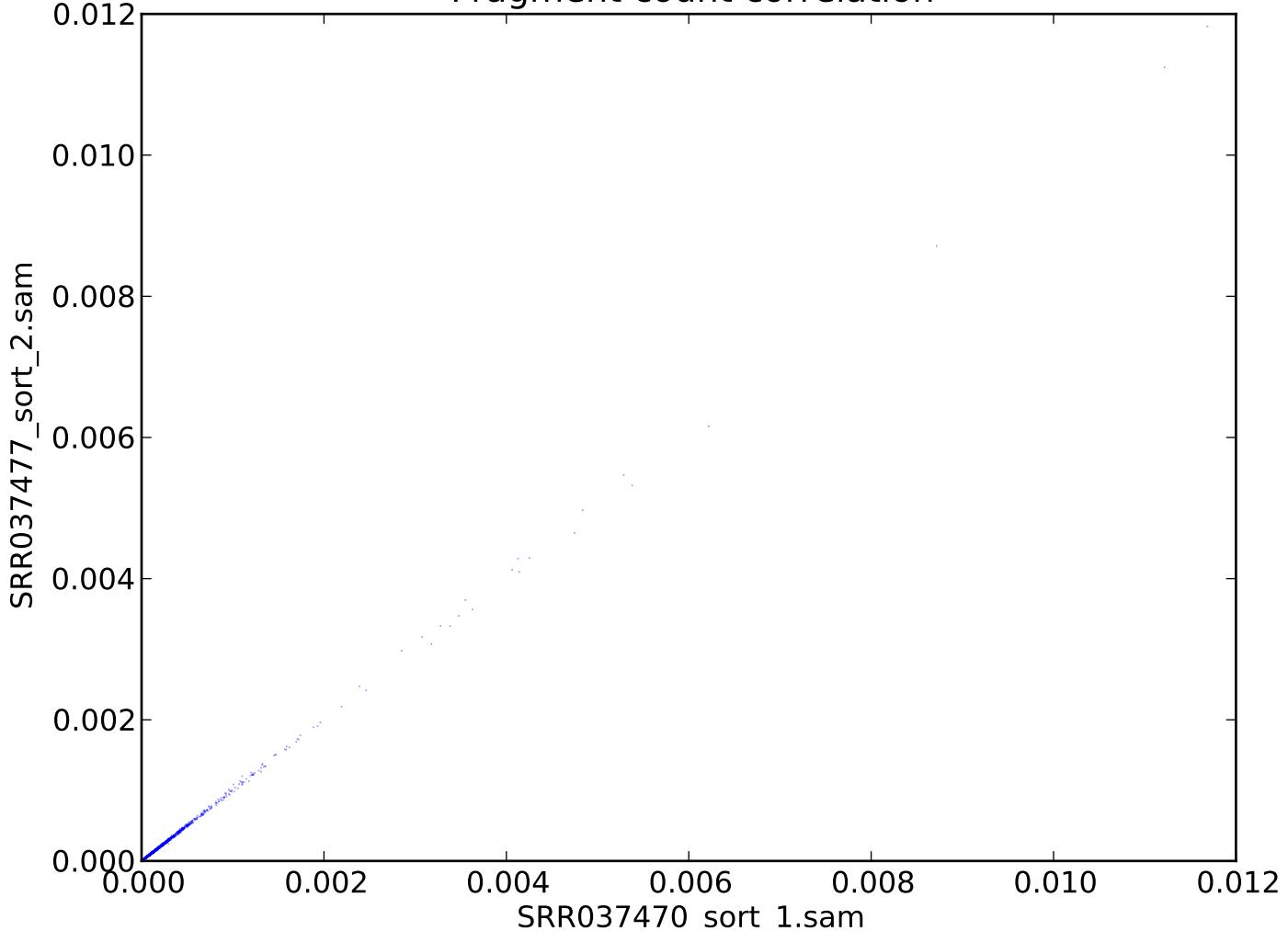
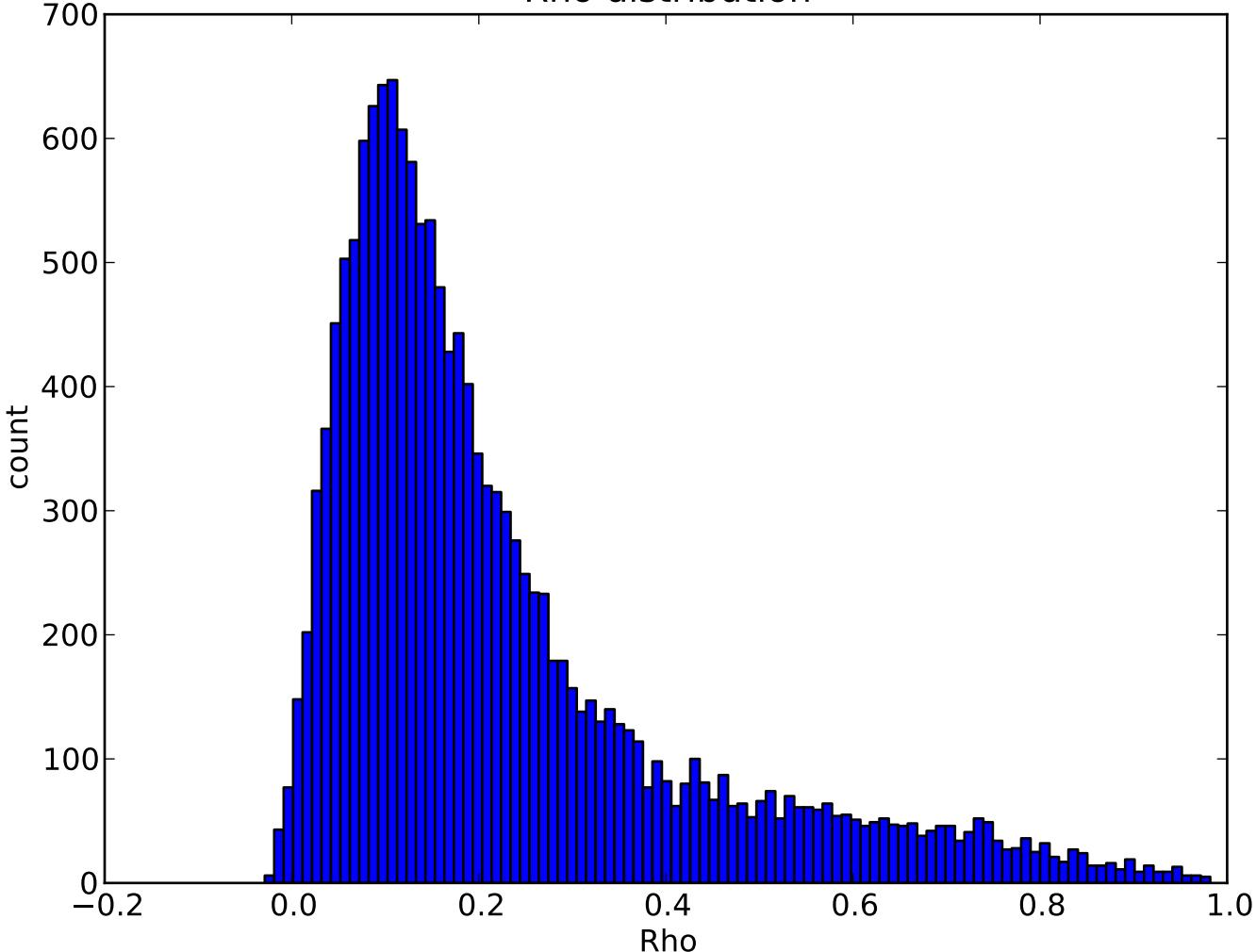


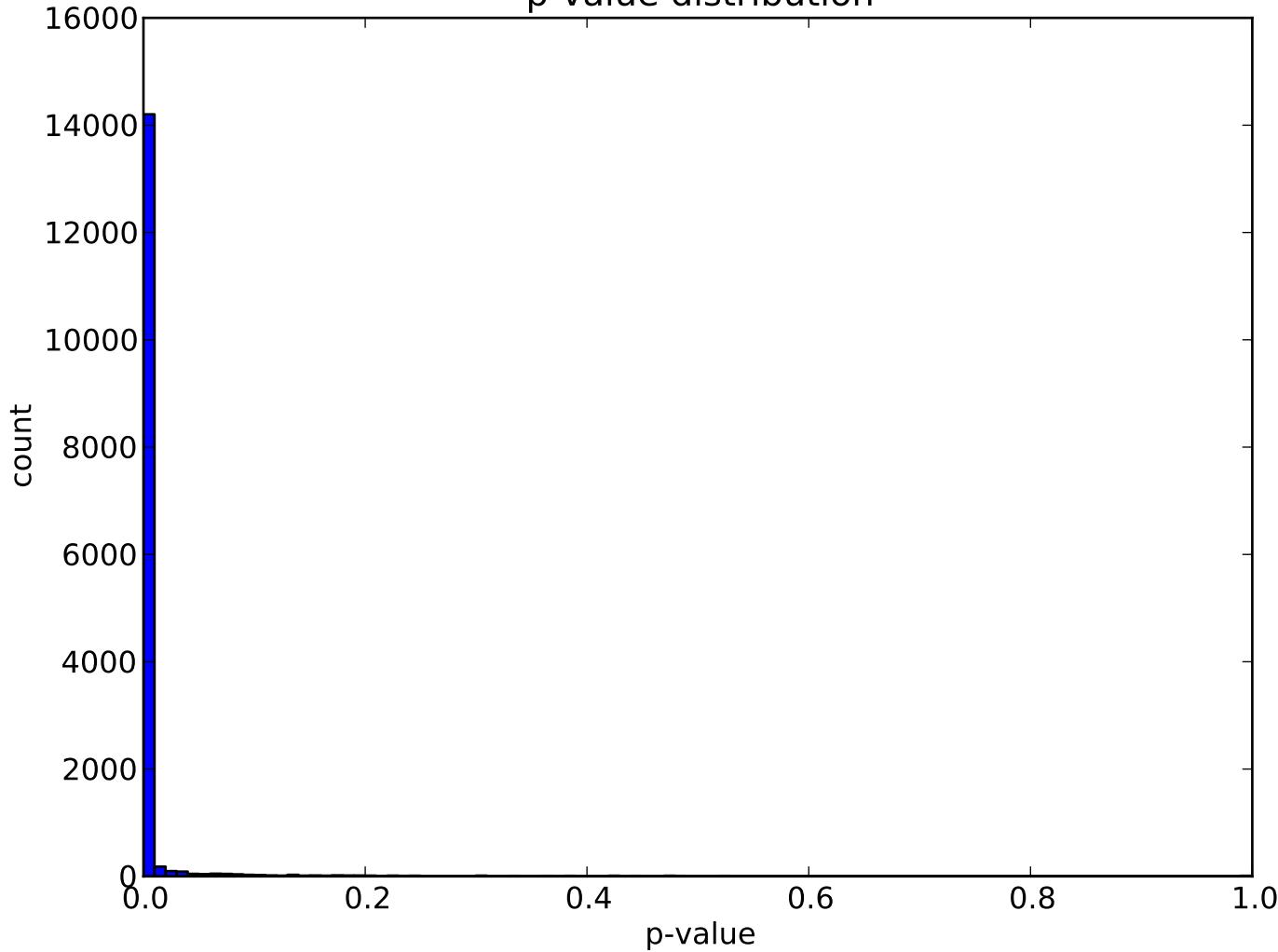
Fragment count correlation



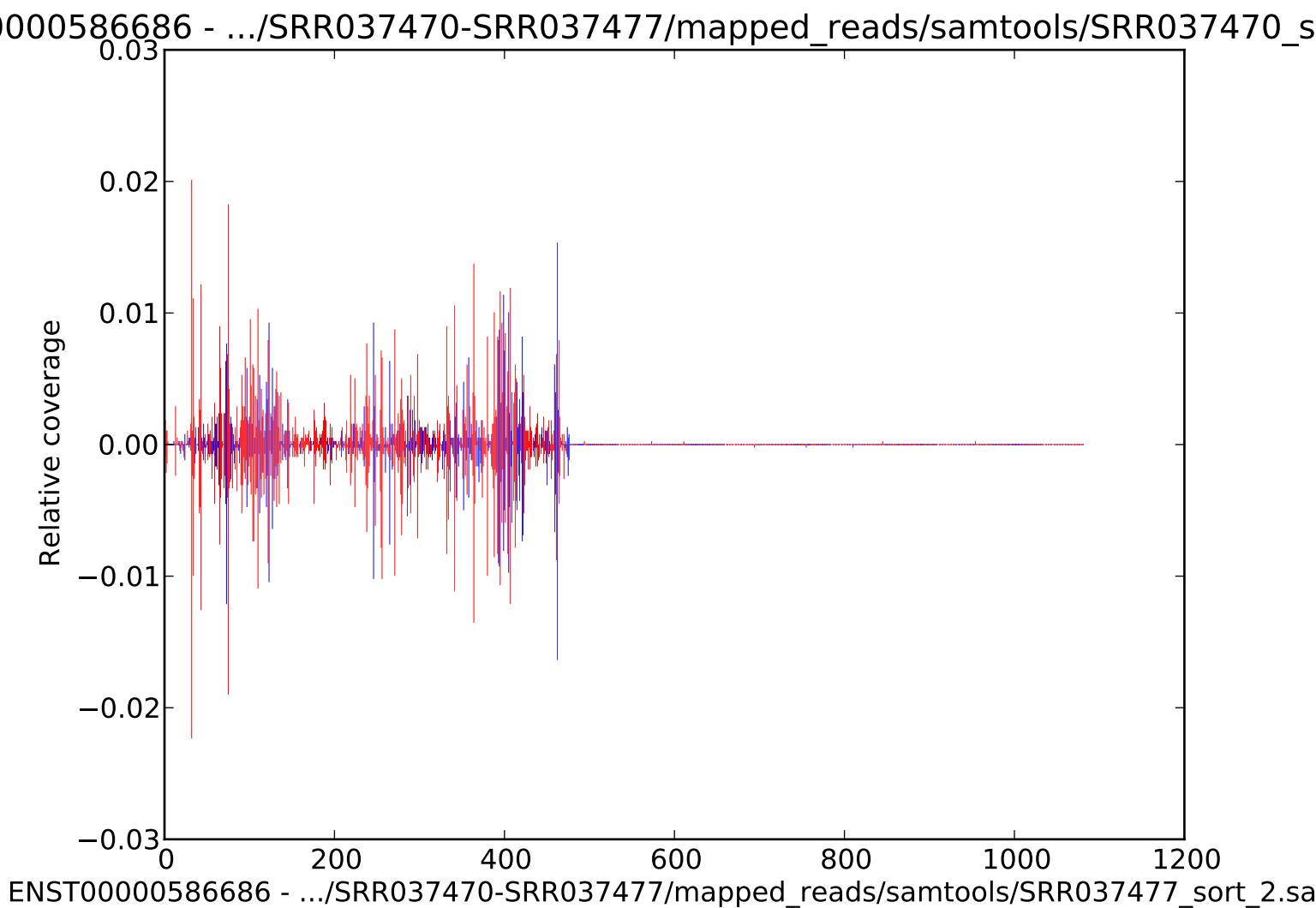
Rho distribution



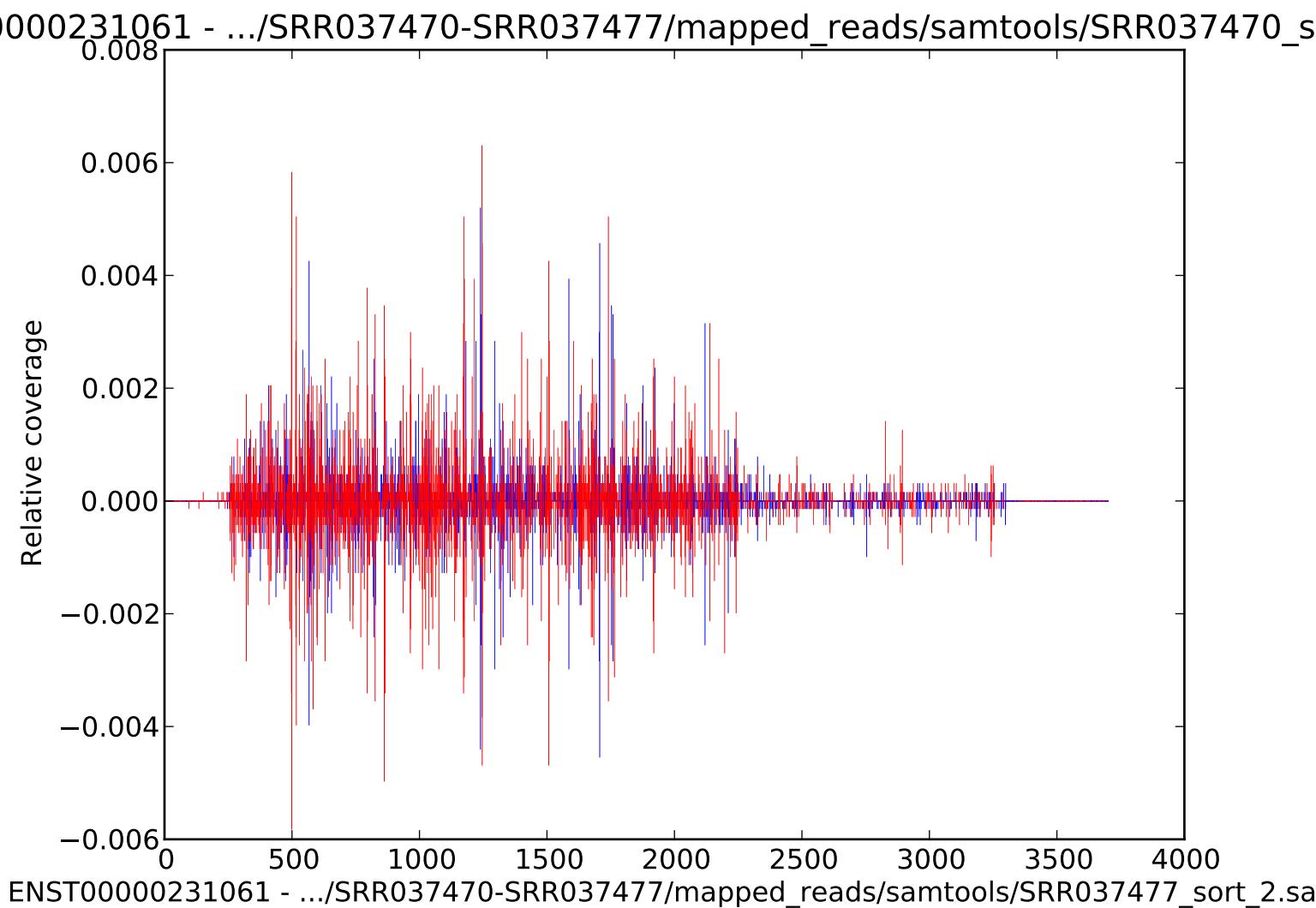
p-value distribution



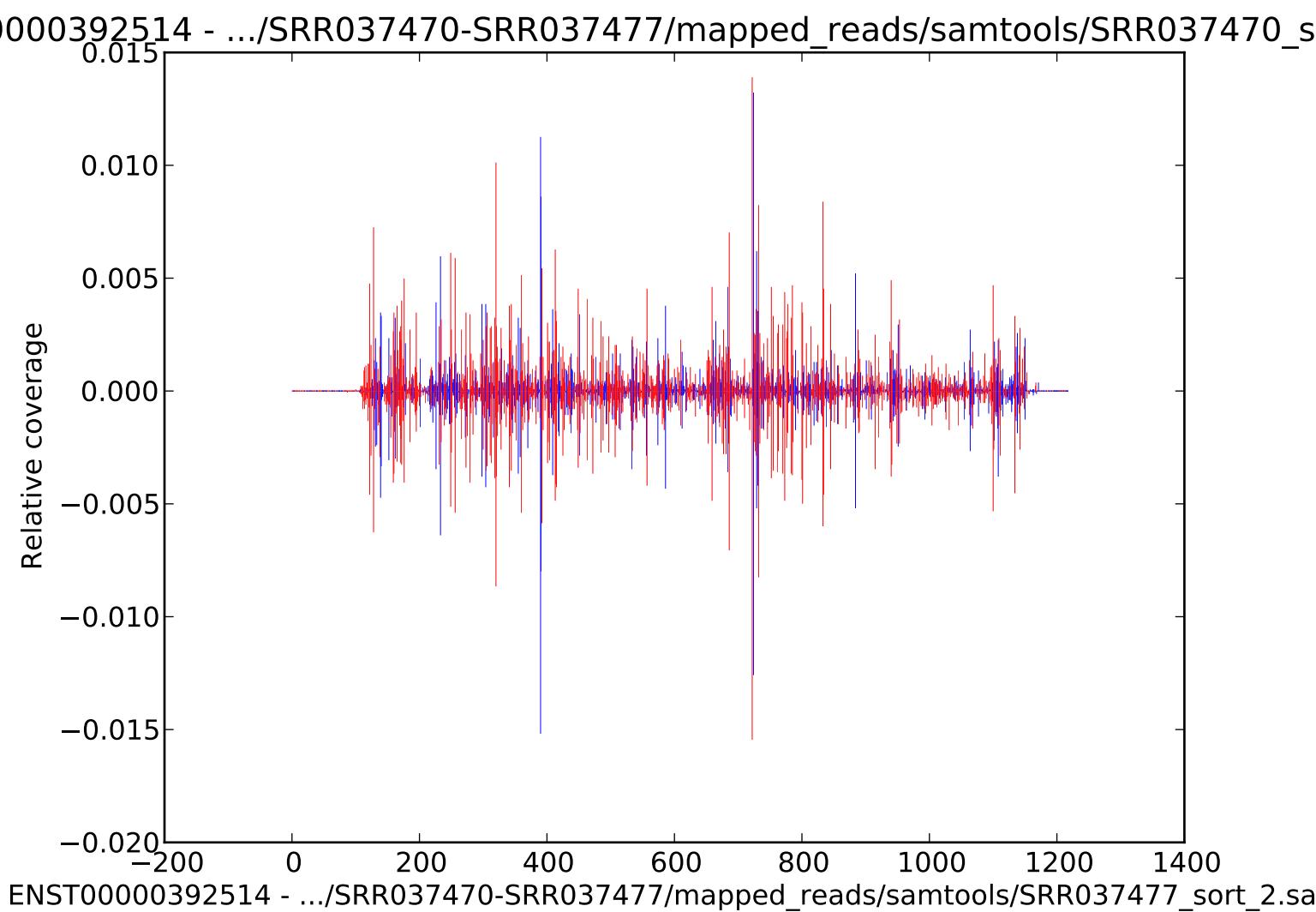
p=0.000000e+00, Rho=9.114407e-01



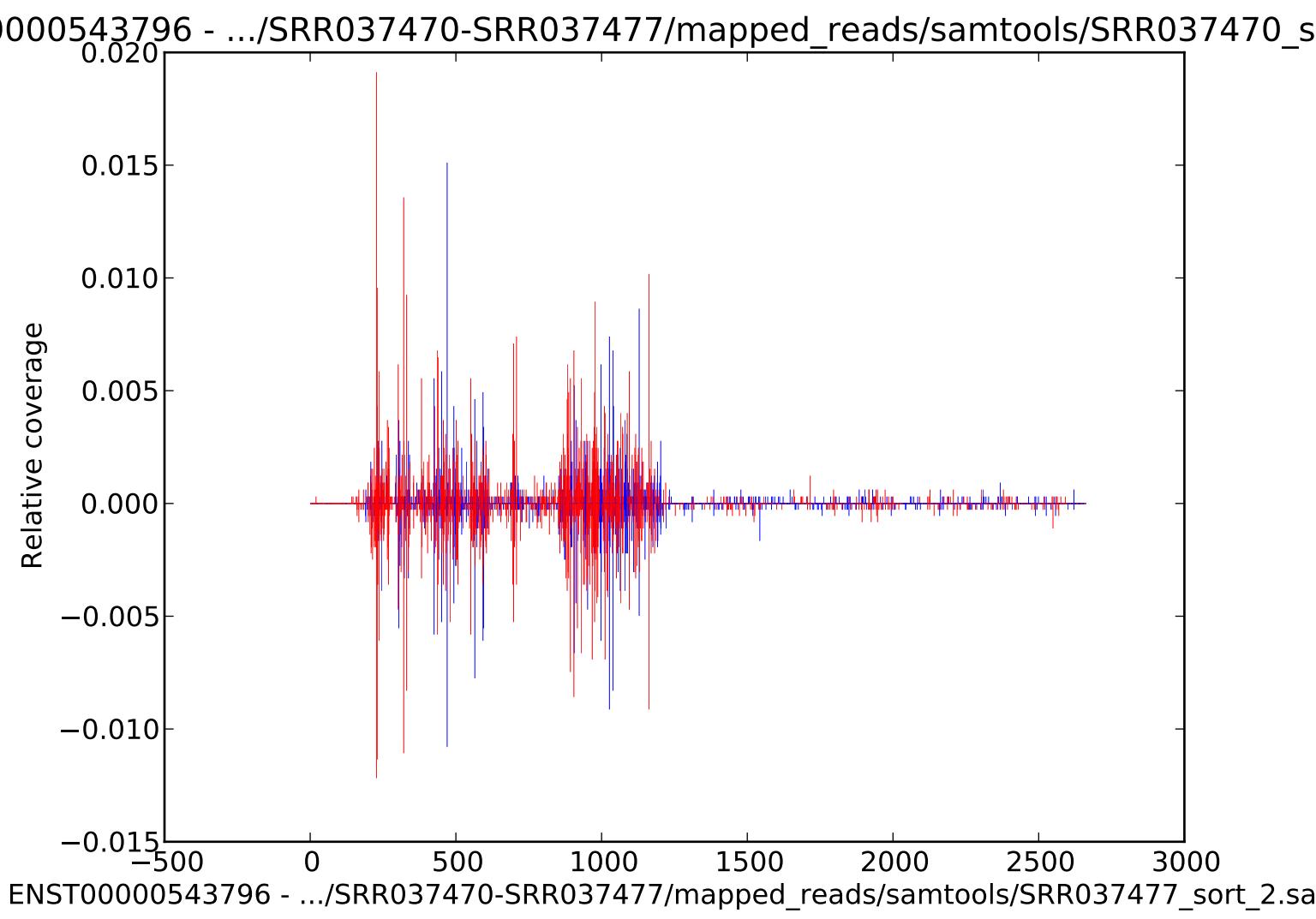
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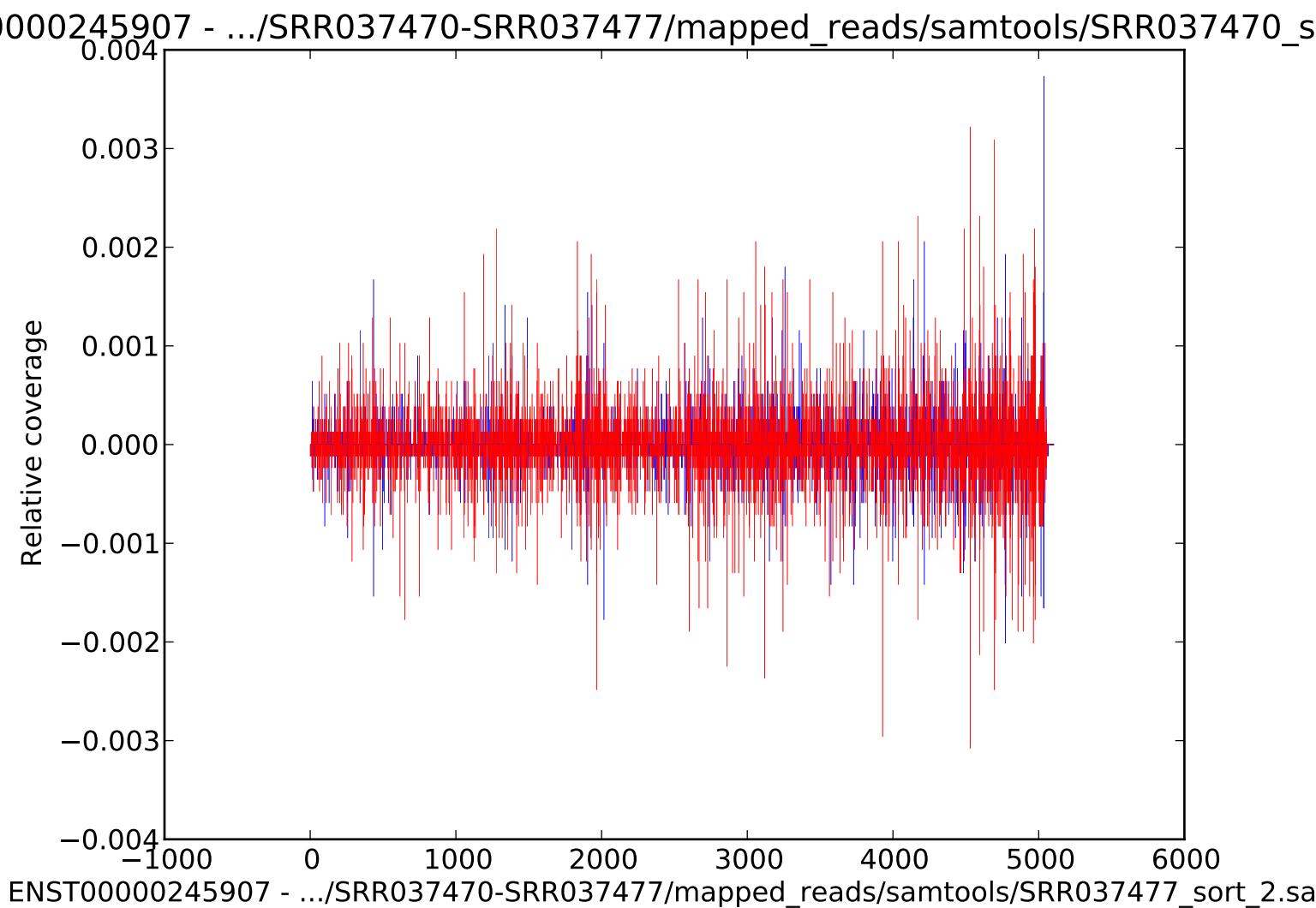
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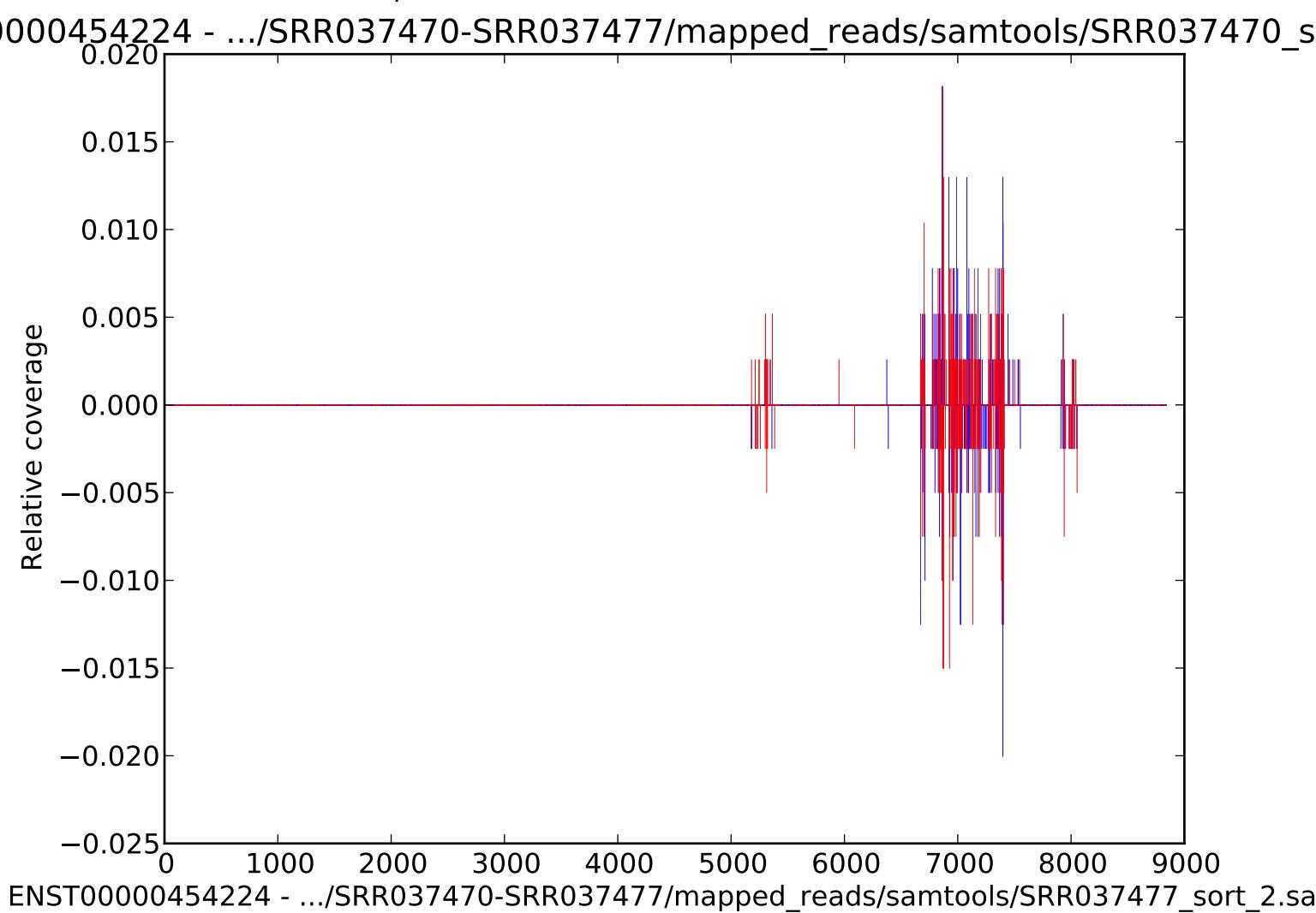
p=0.000000e+00, Rho=6.640880e-01



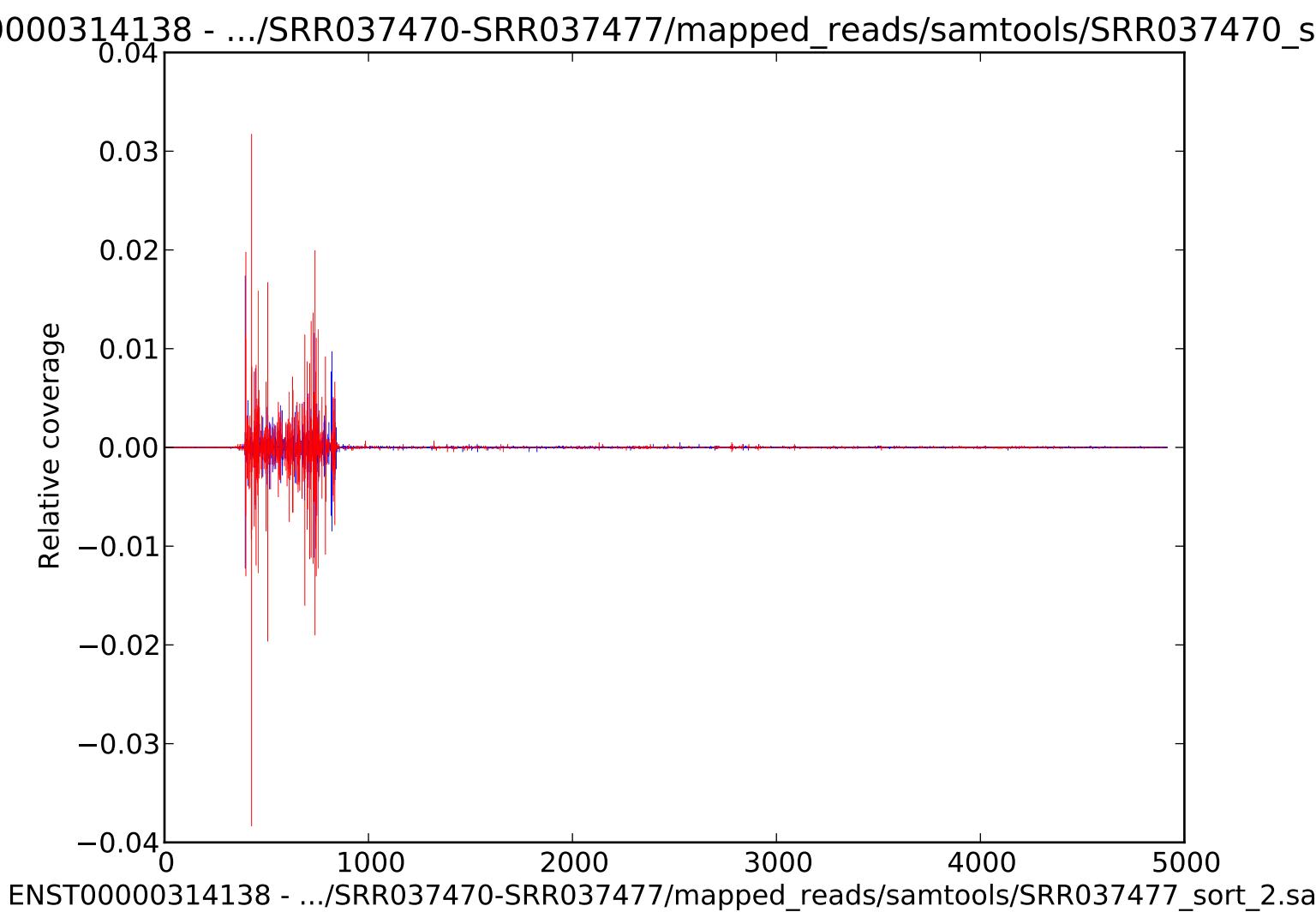
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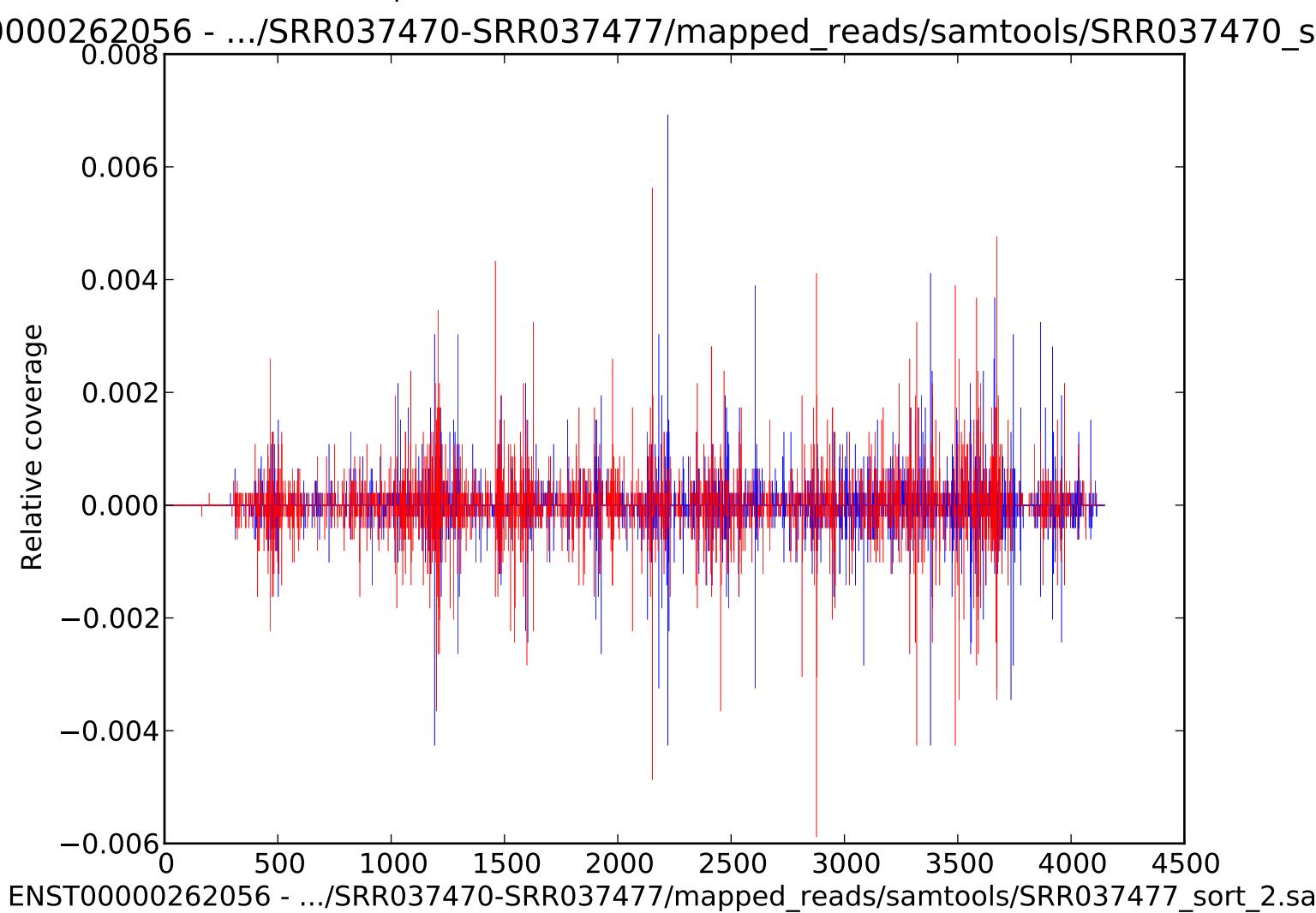
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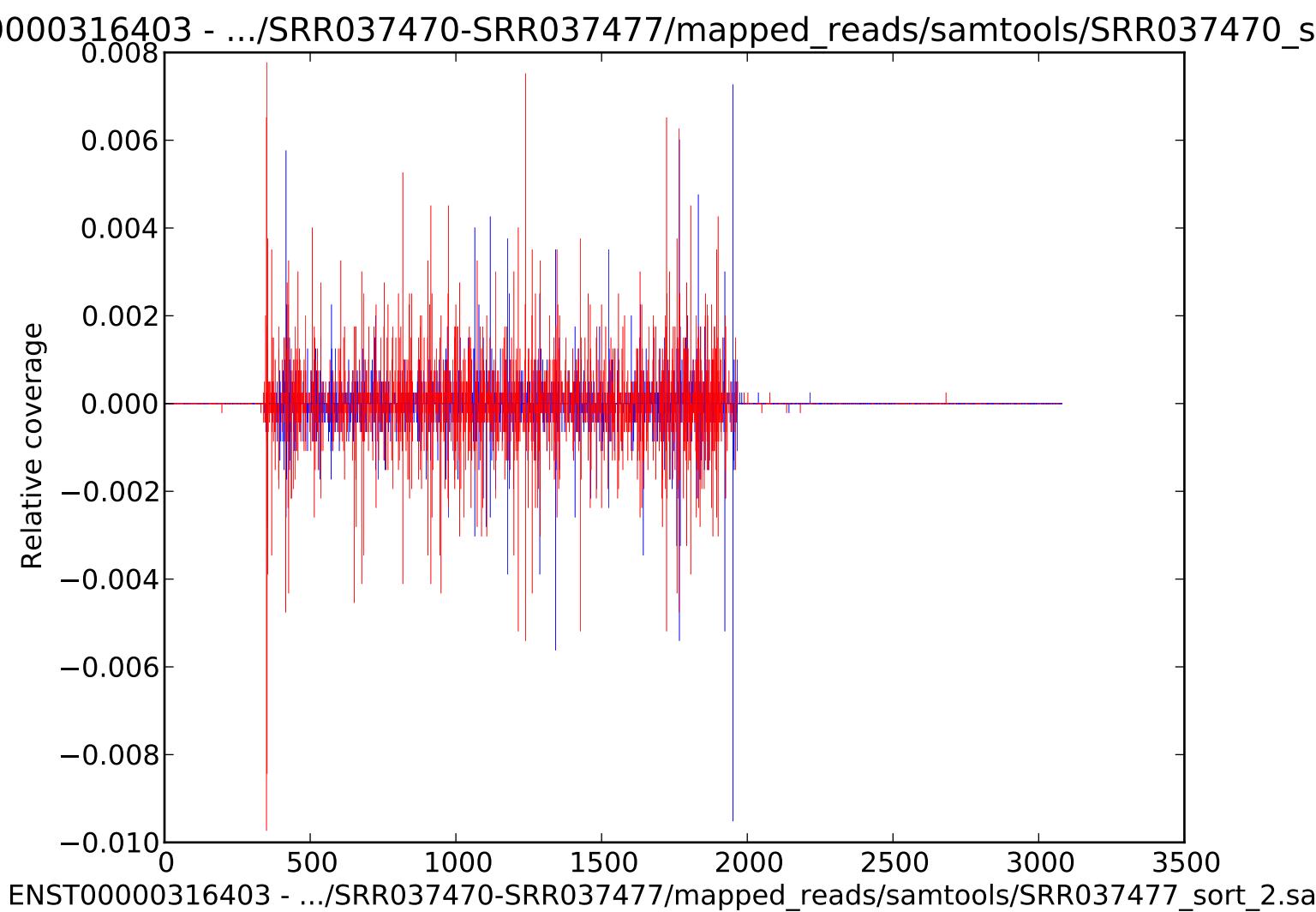
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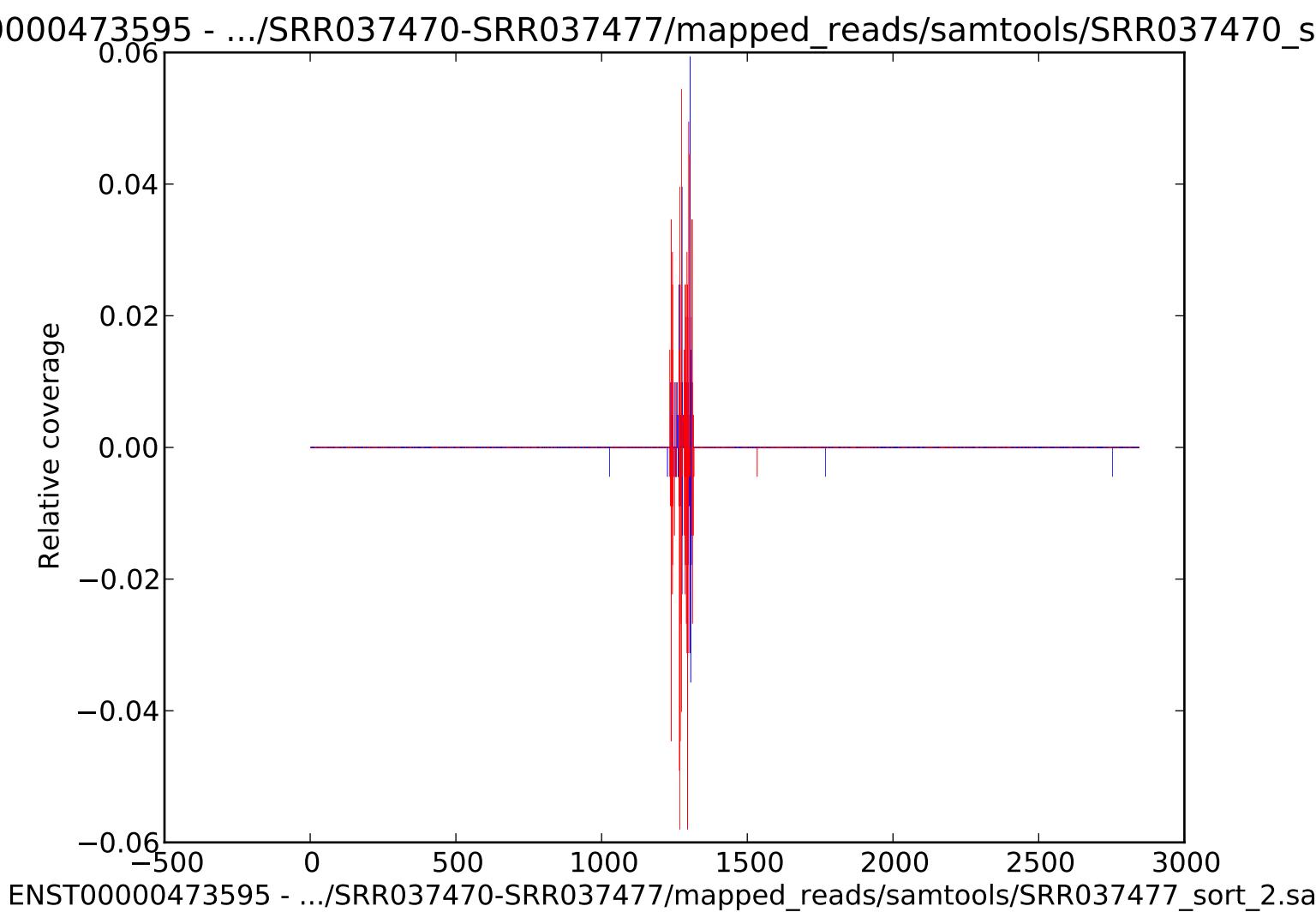
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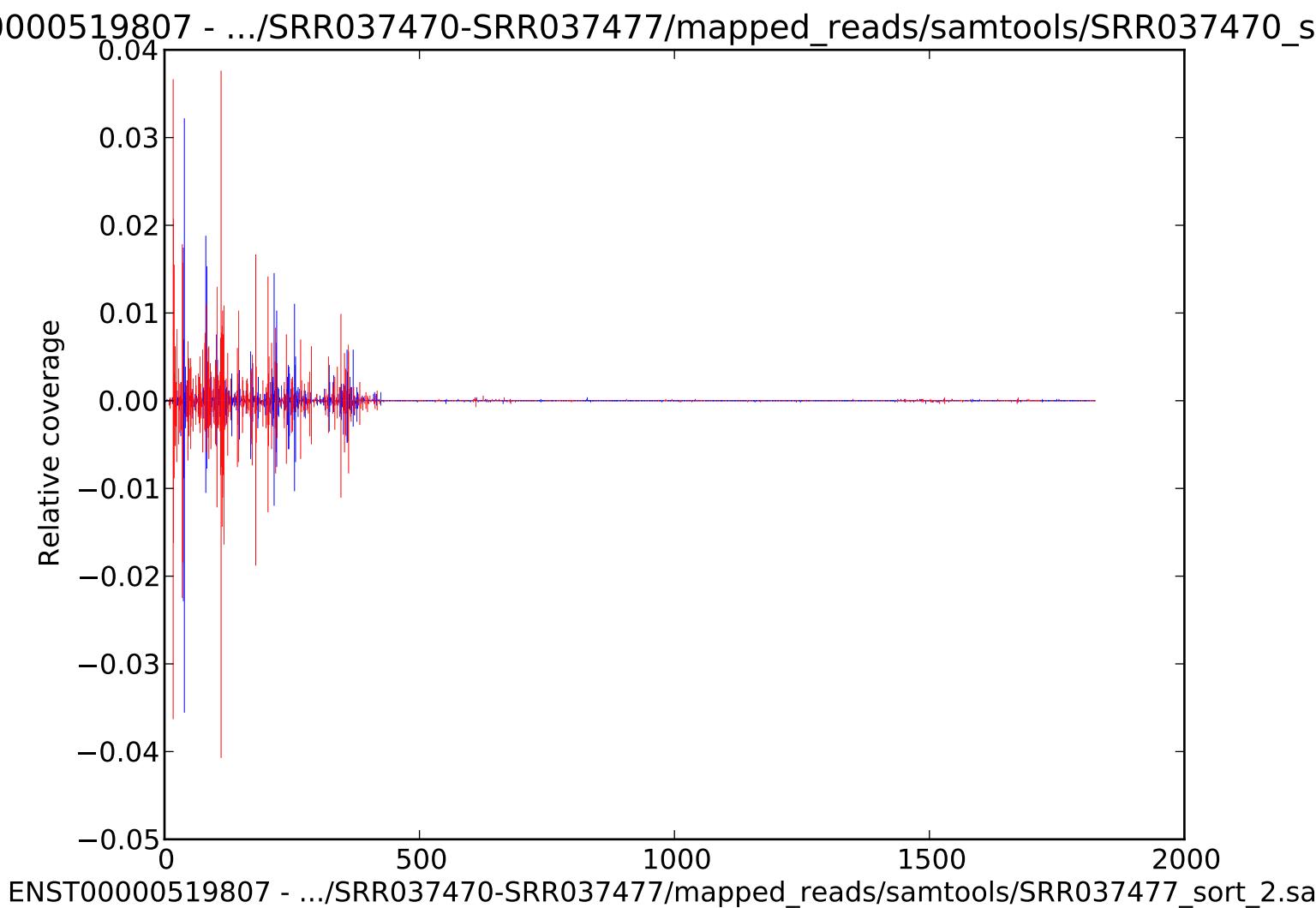
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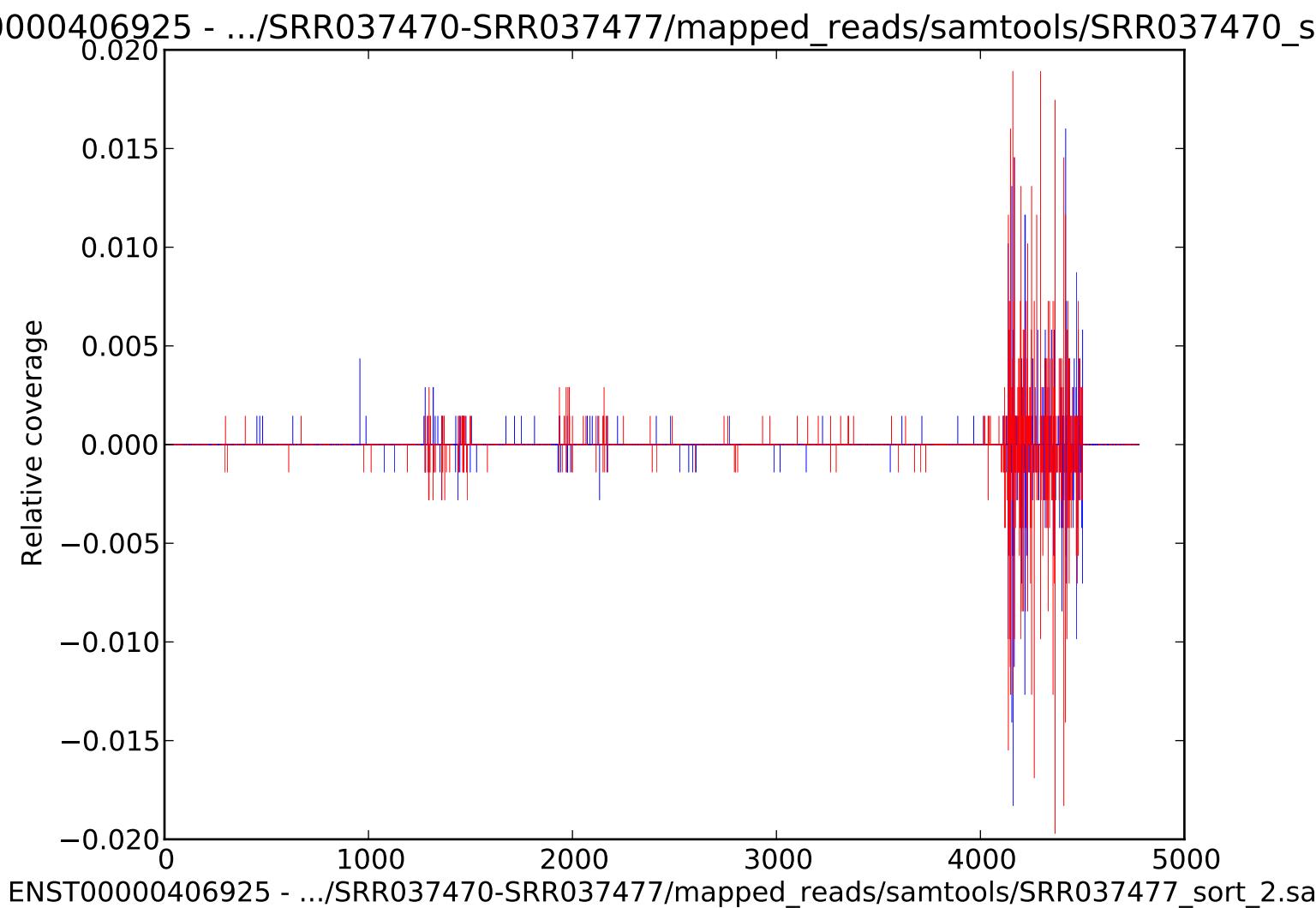
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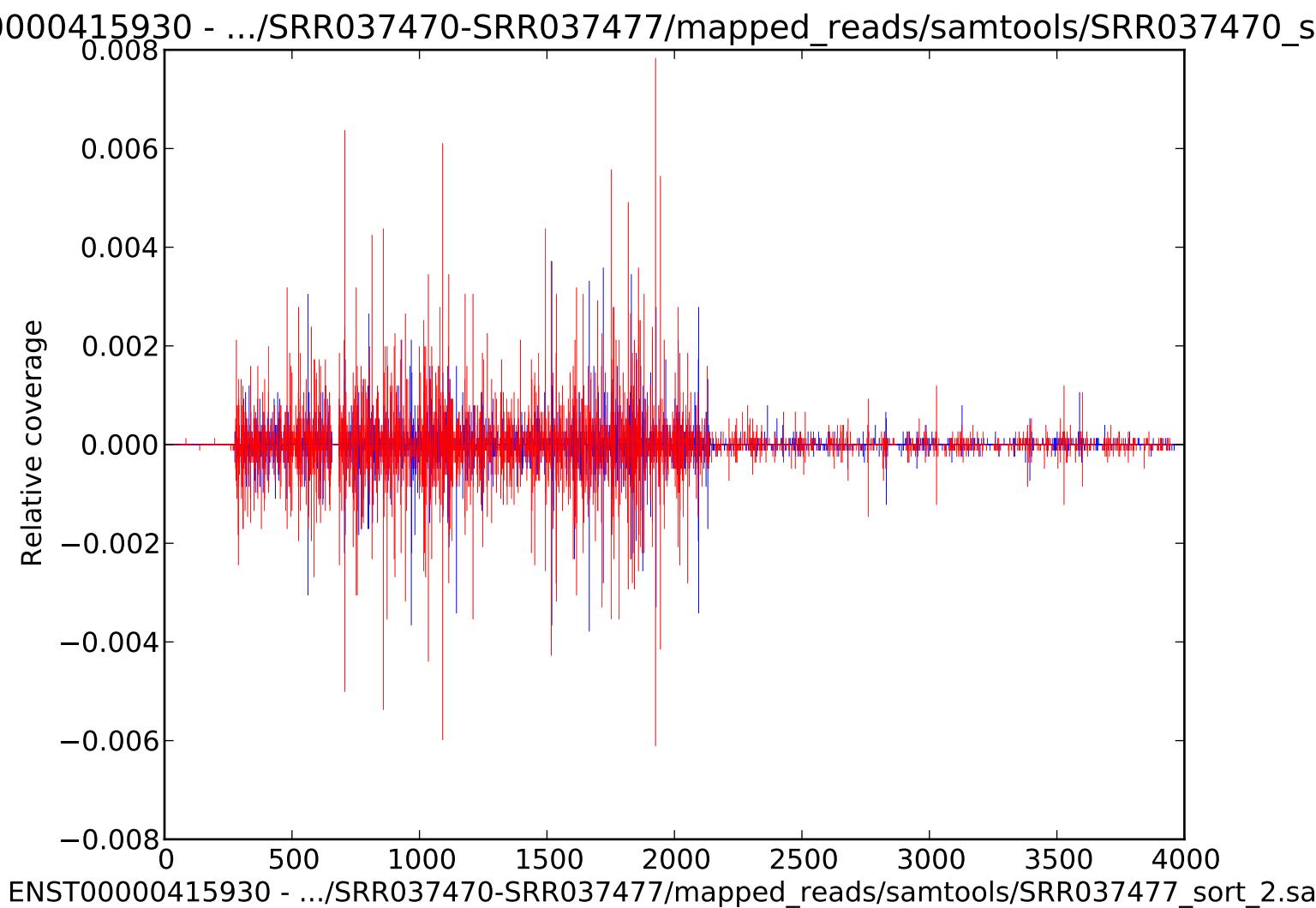
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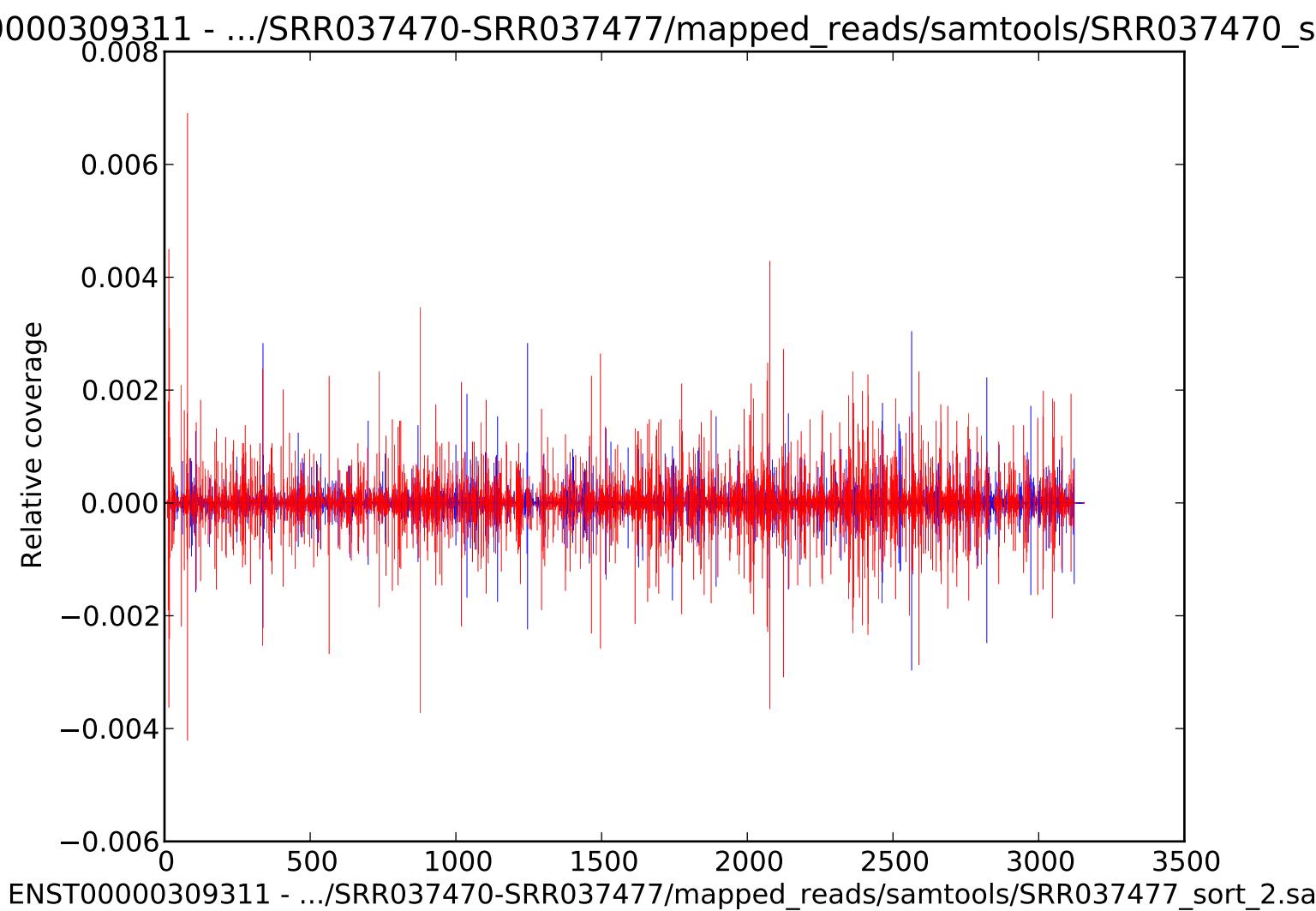
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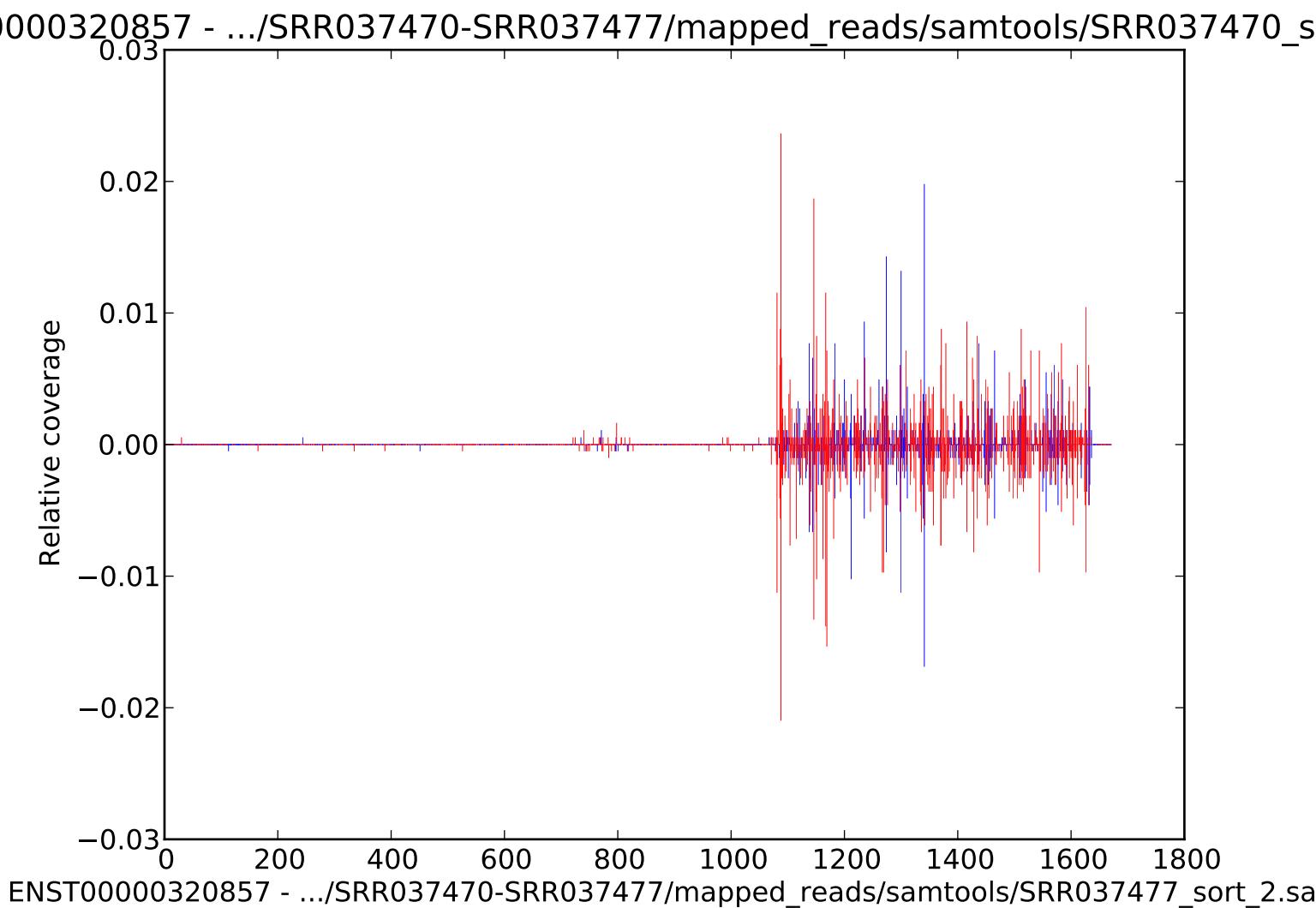
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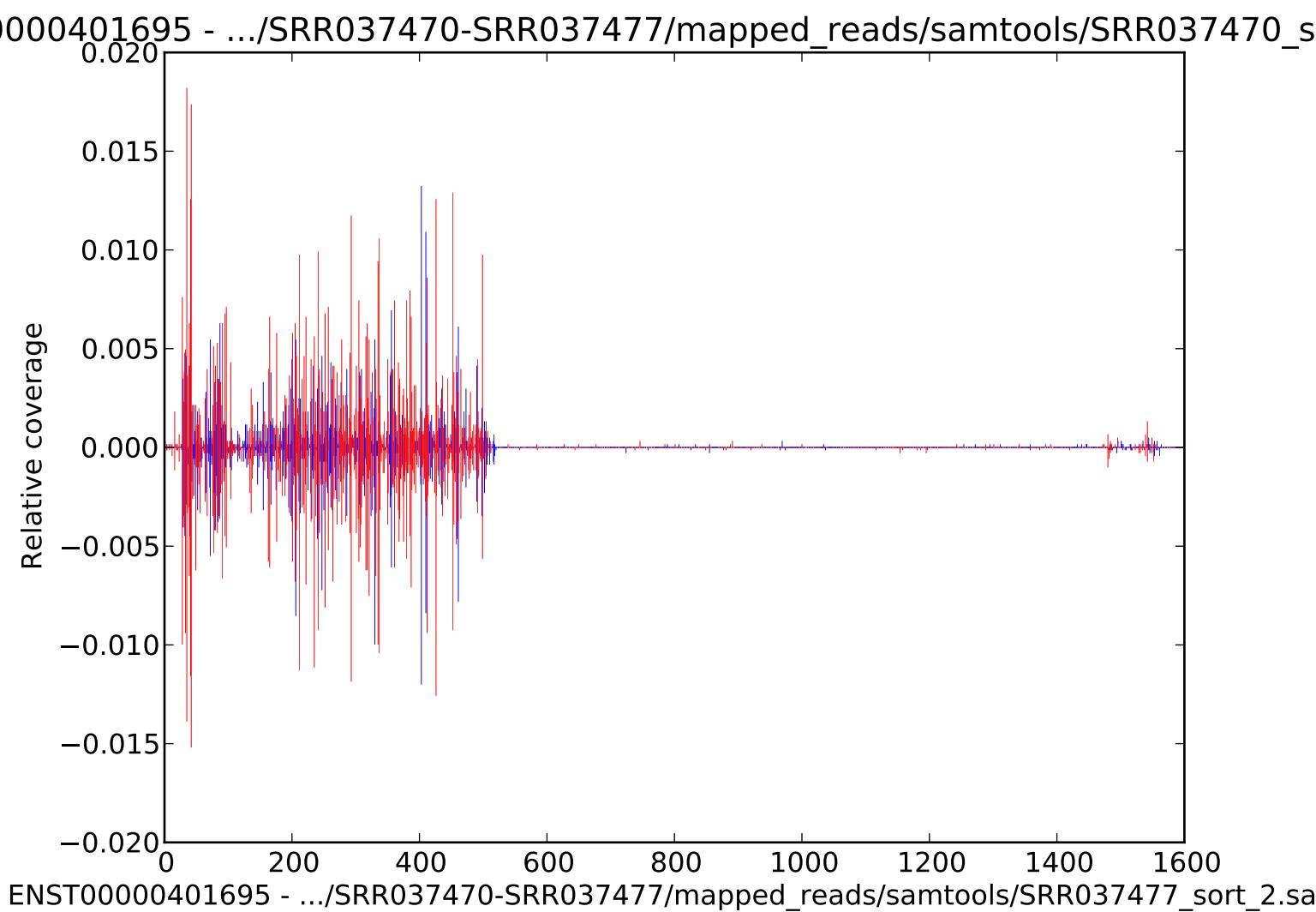
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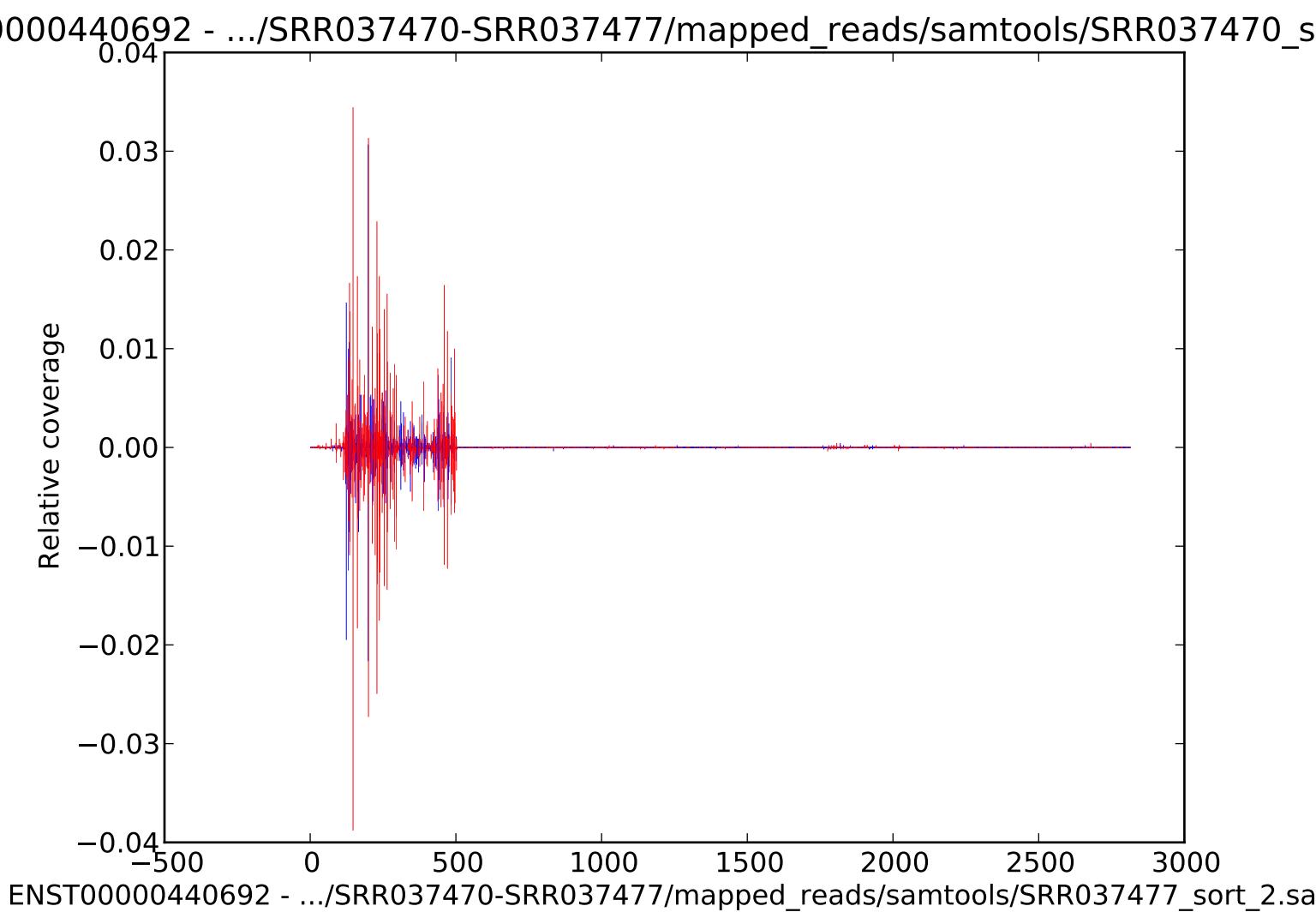
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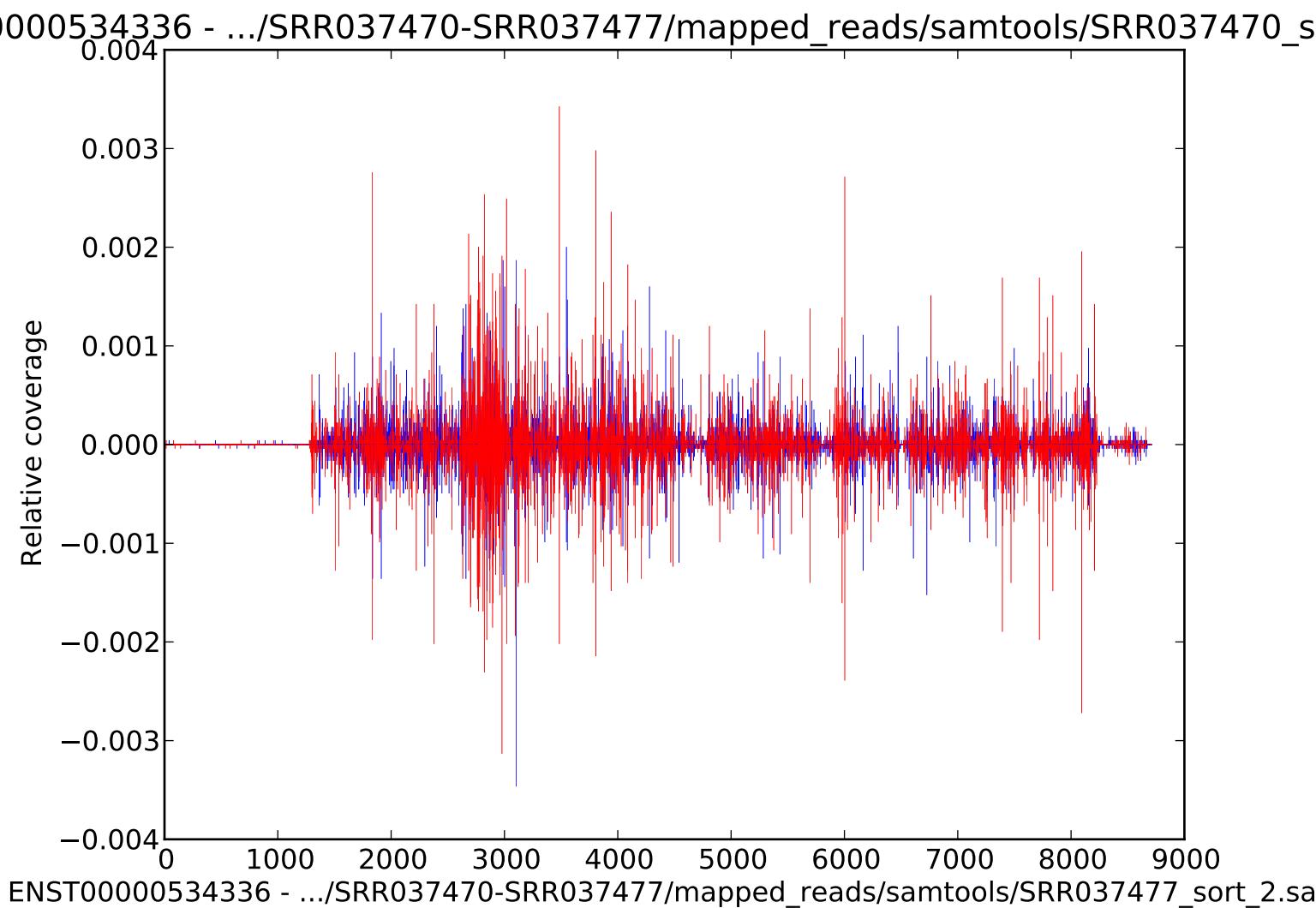
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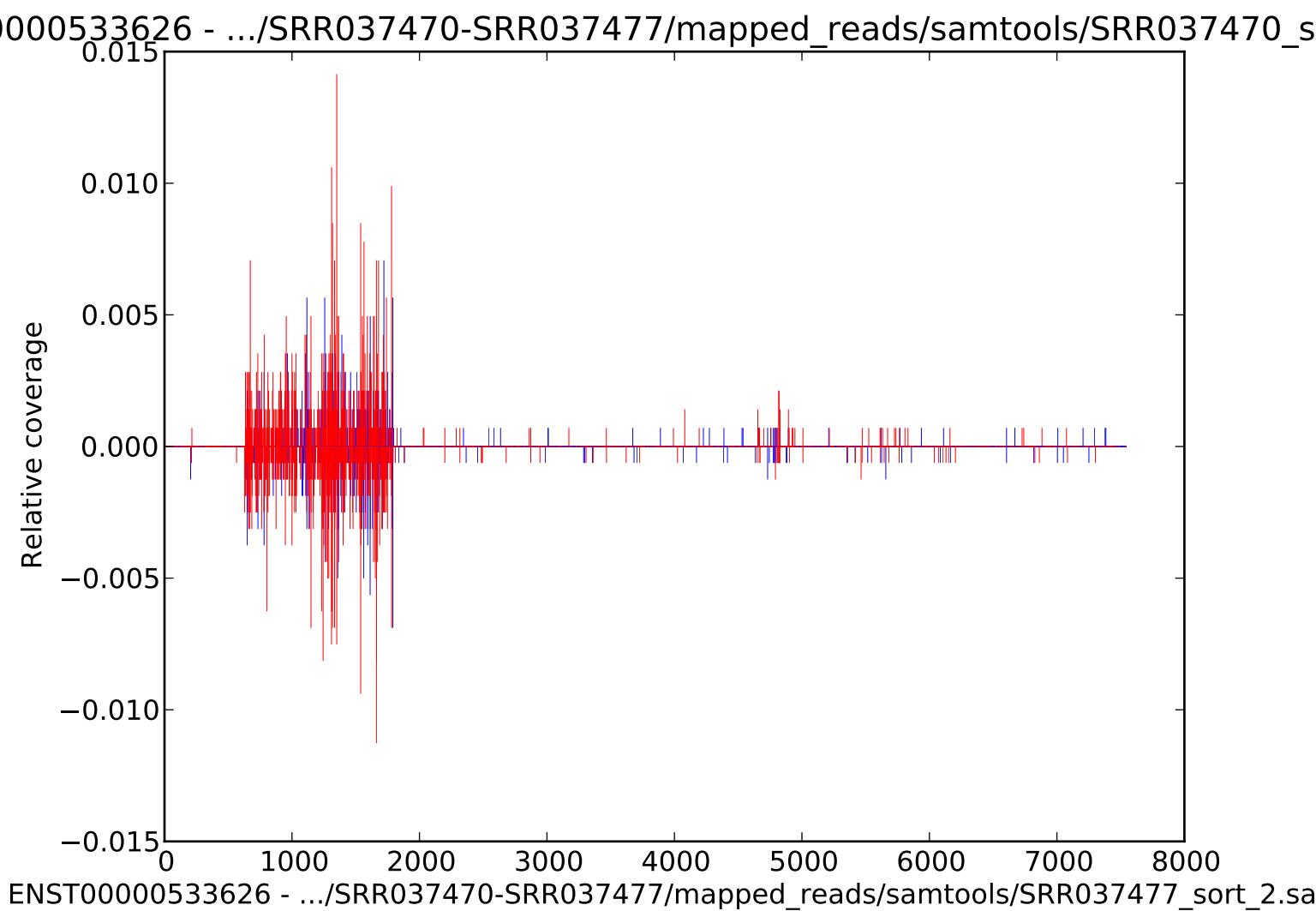
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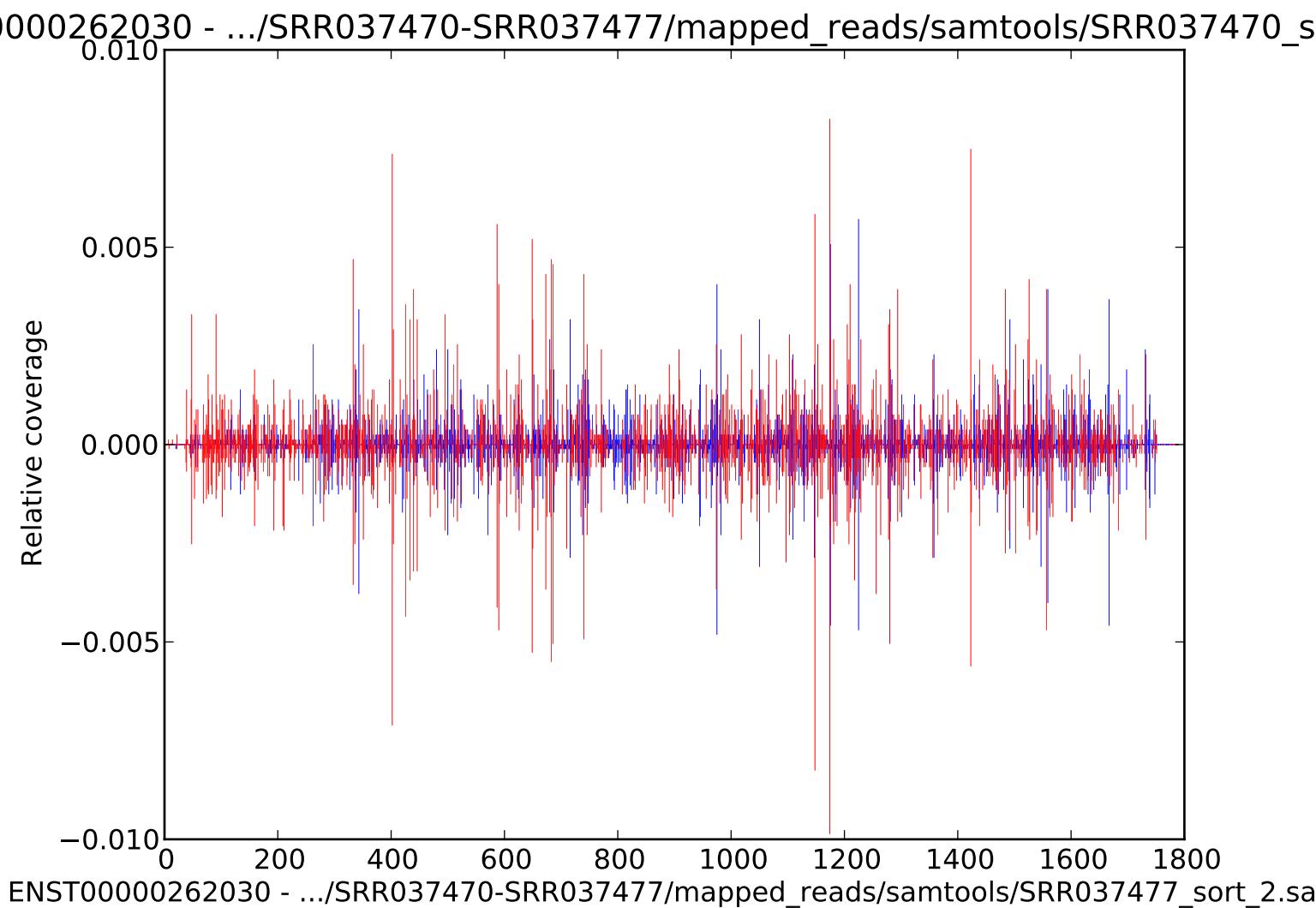
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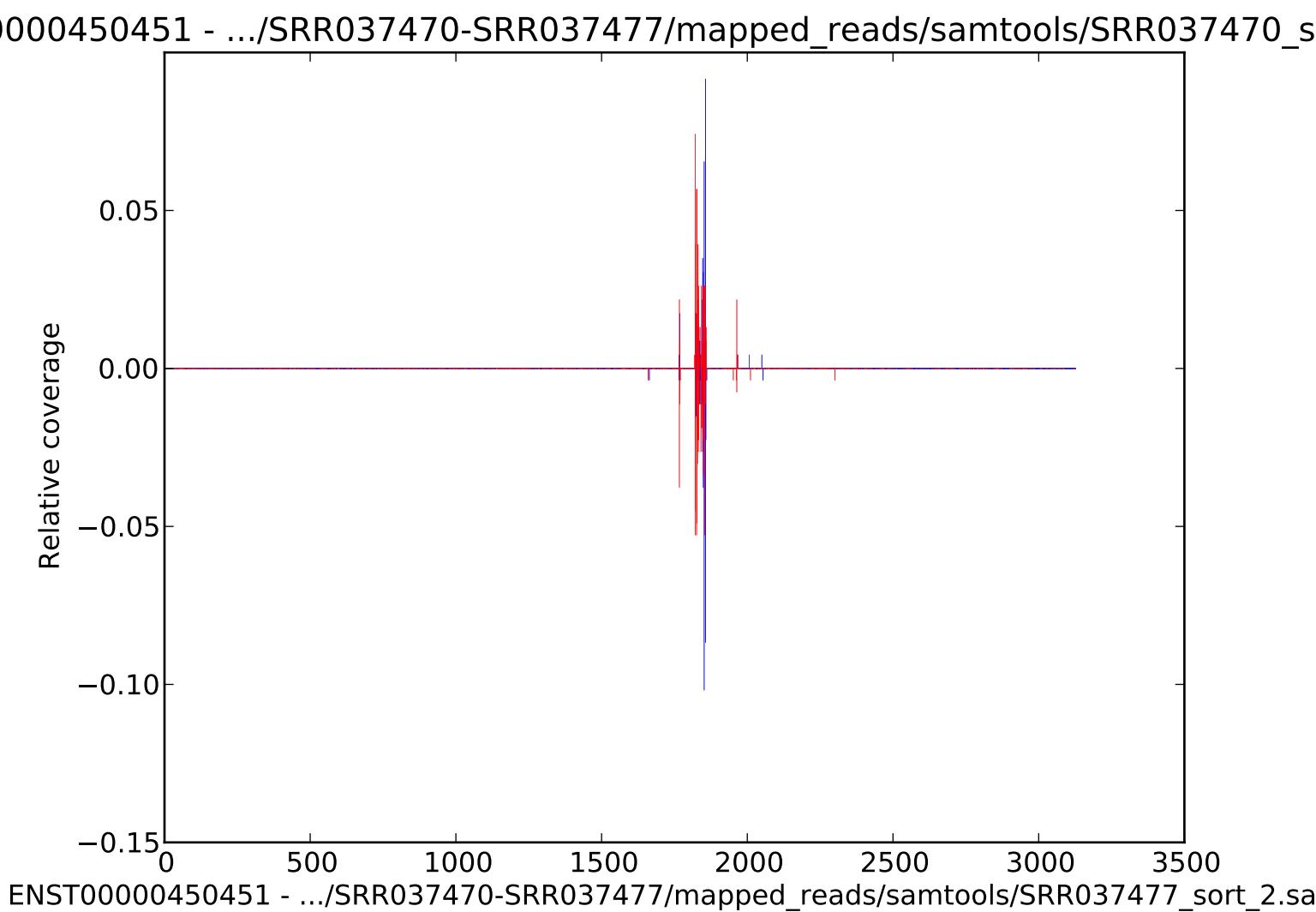
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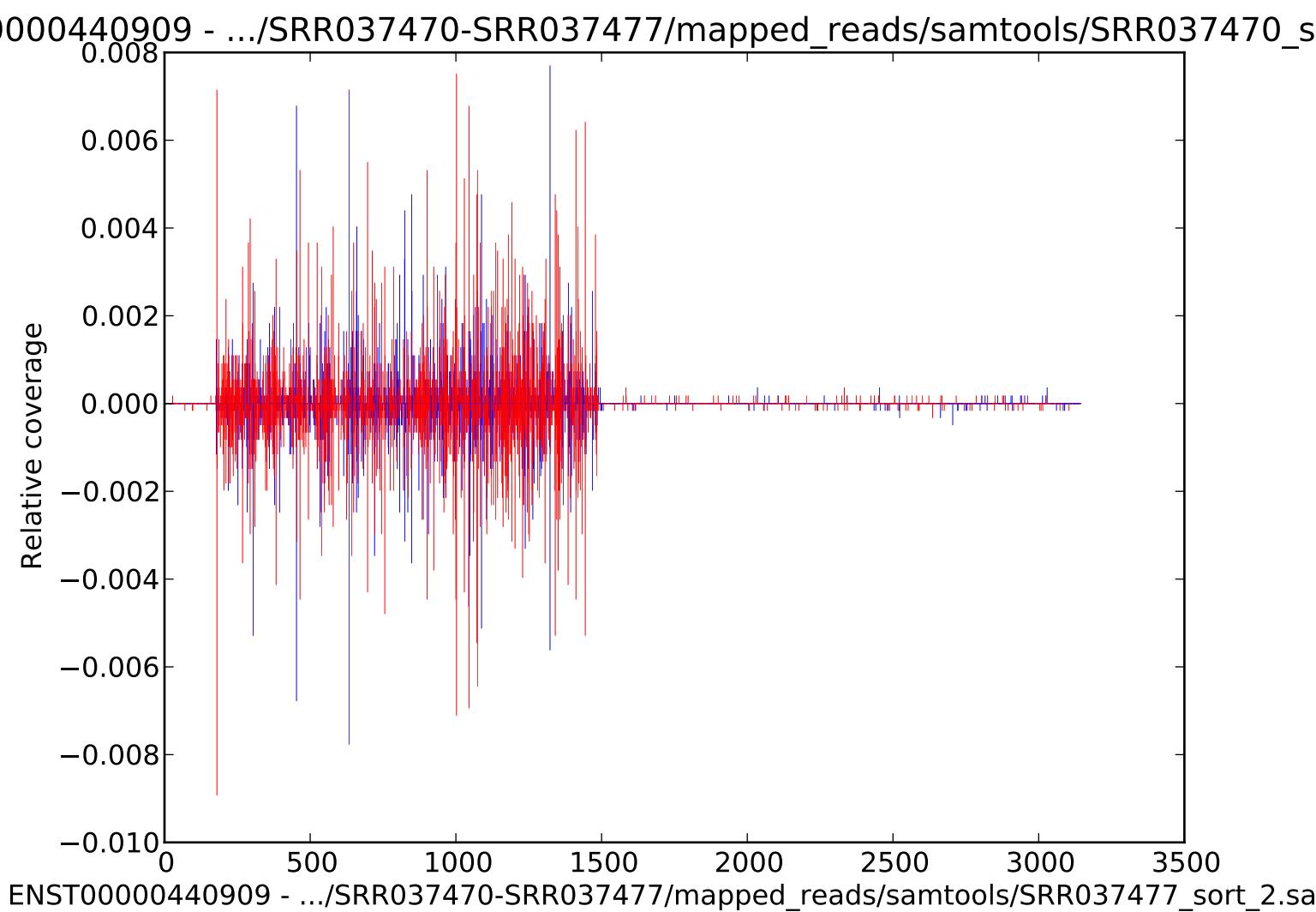
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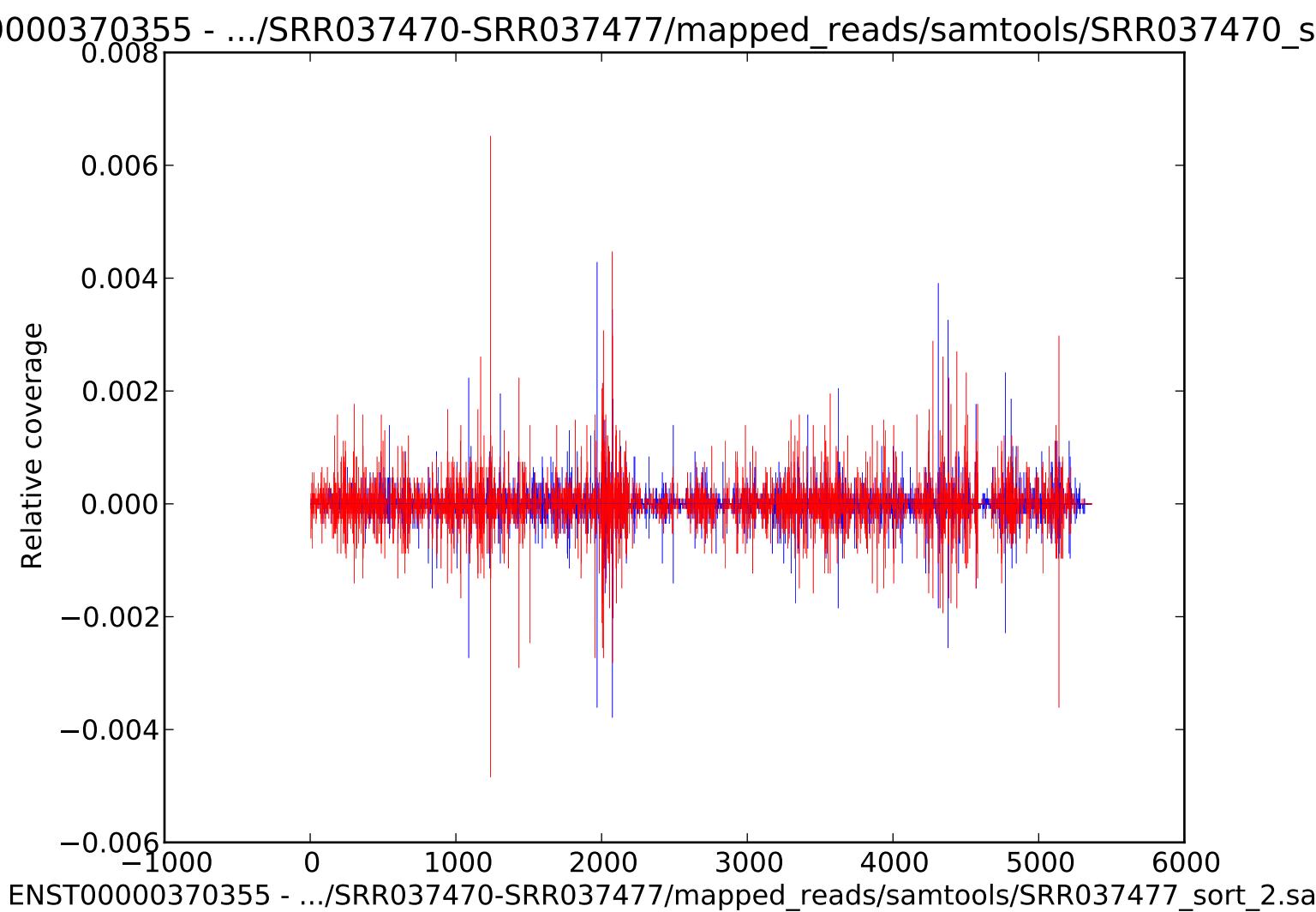
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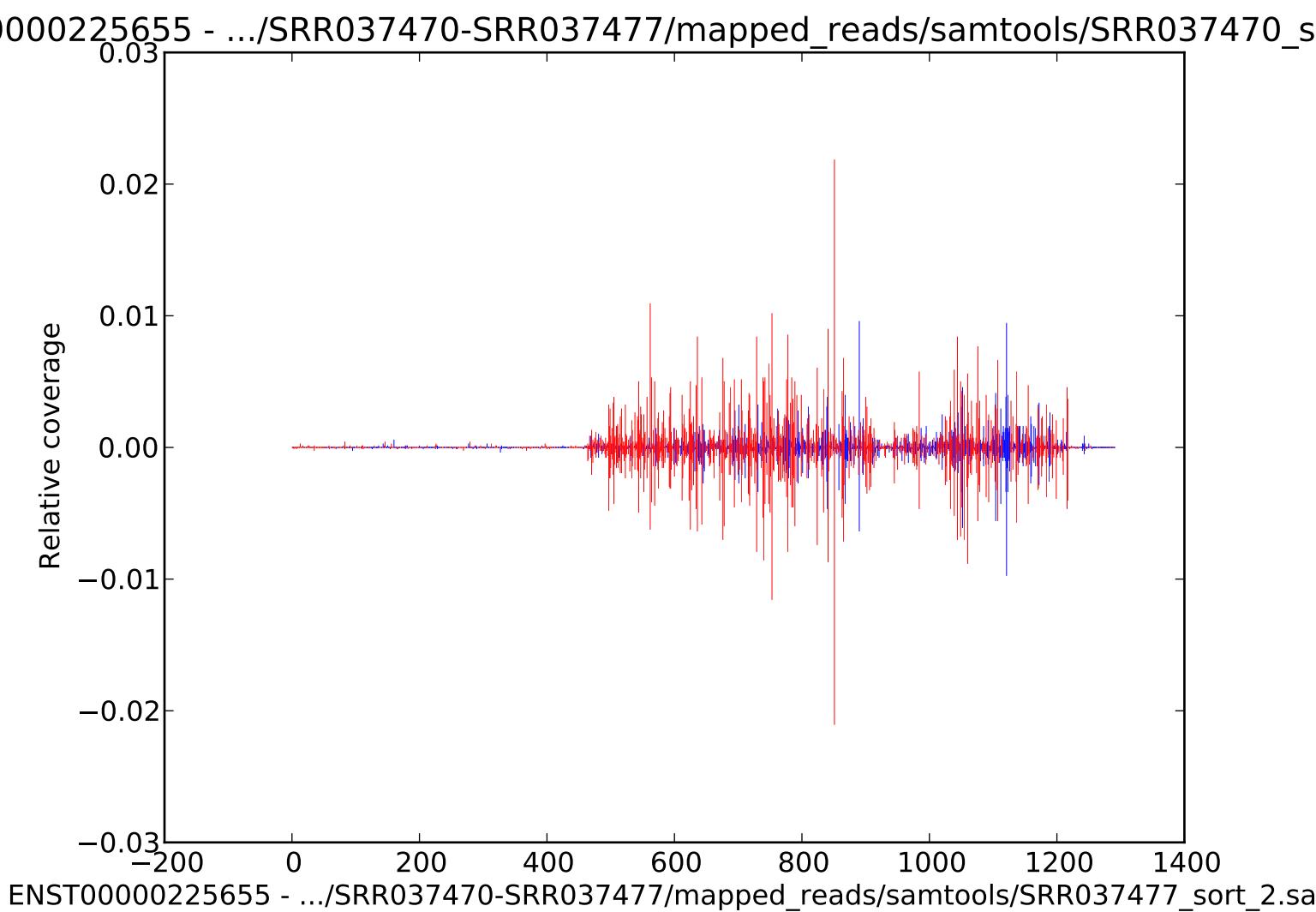
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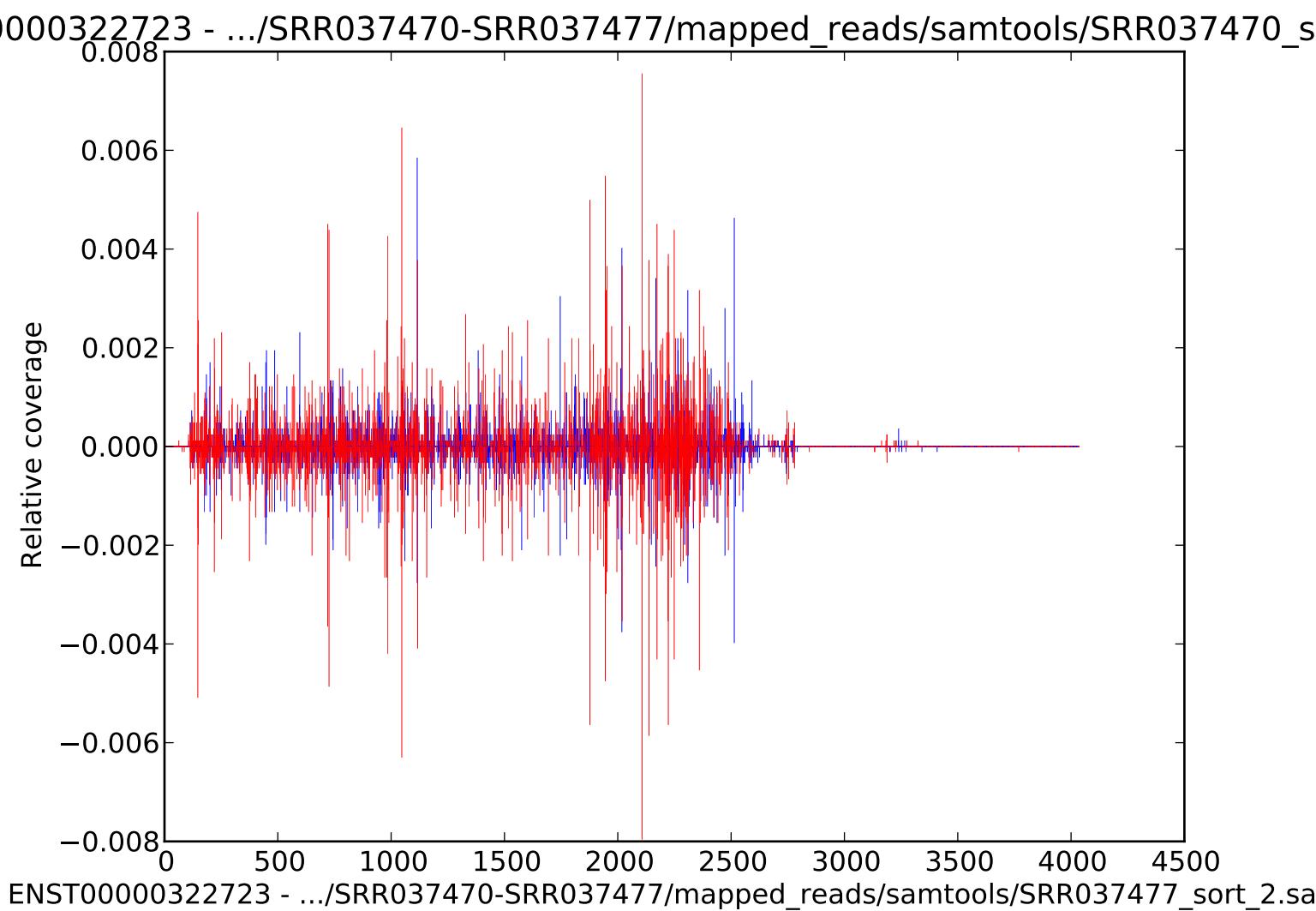
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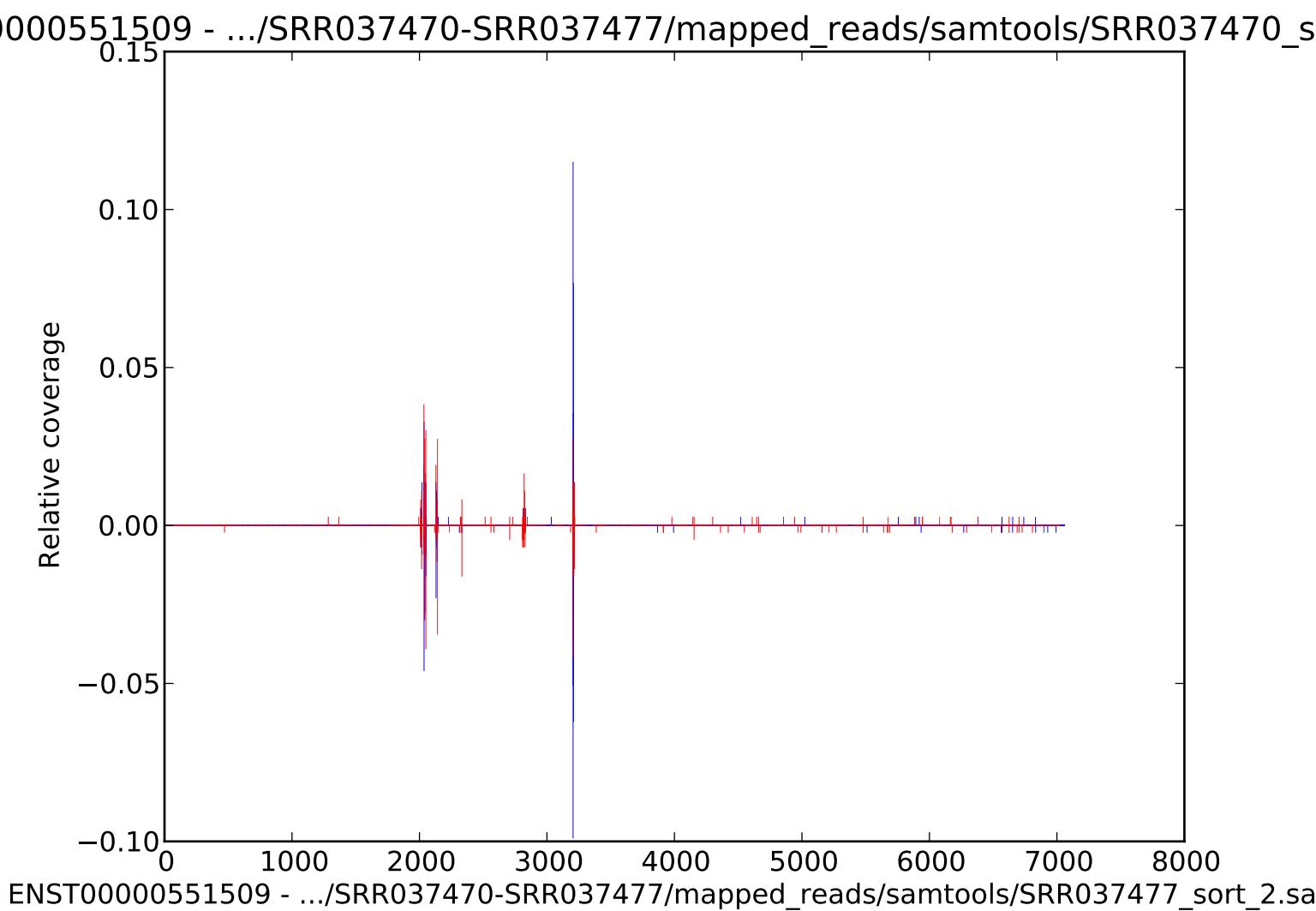
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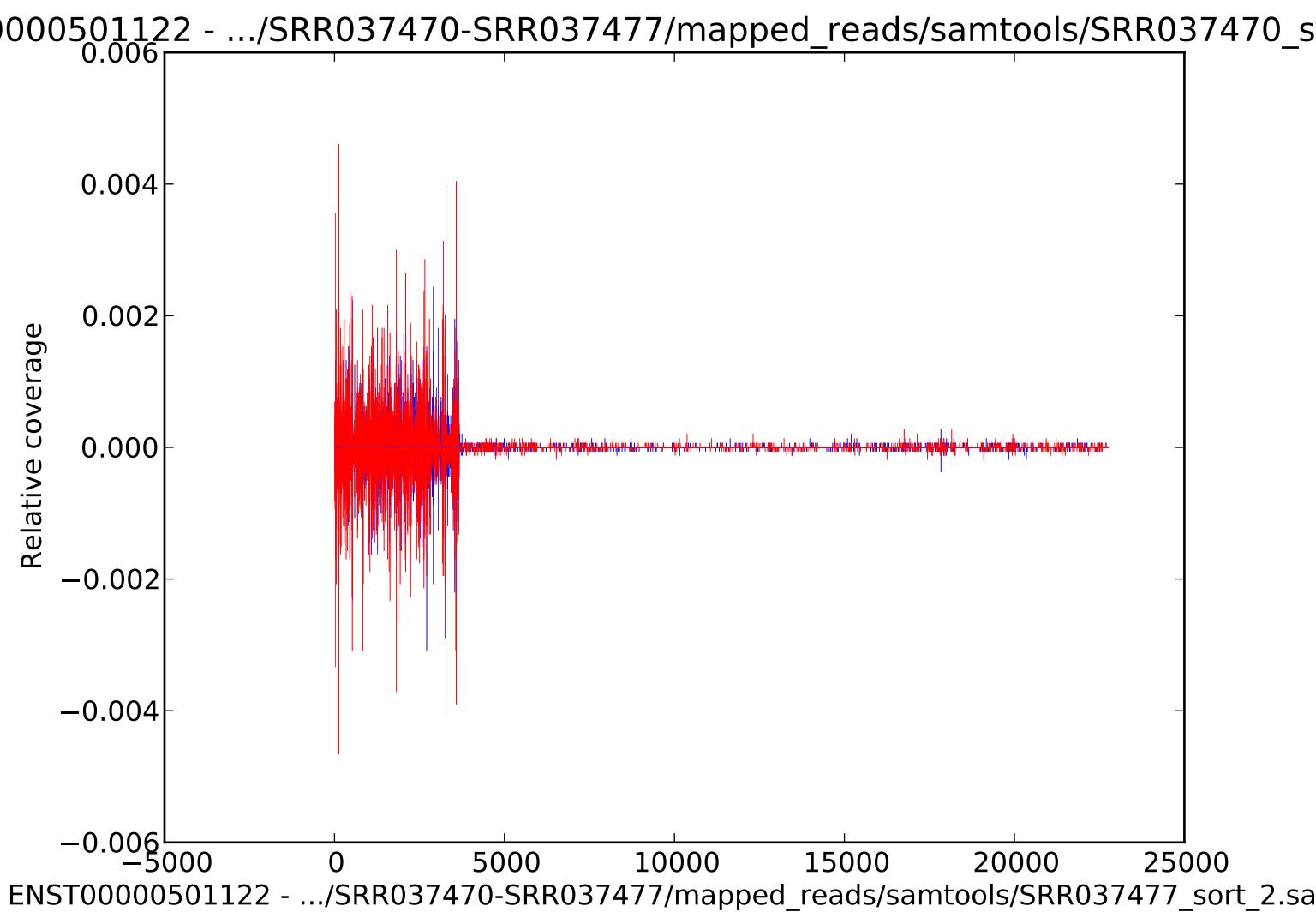
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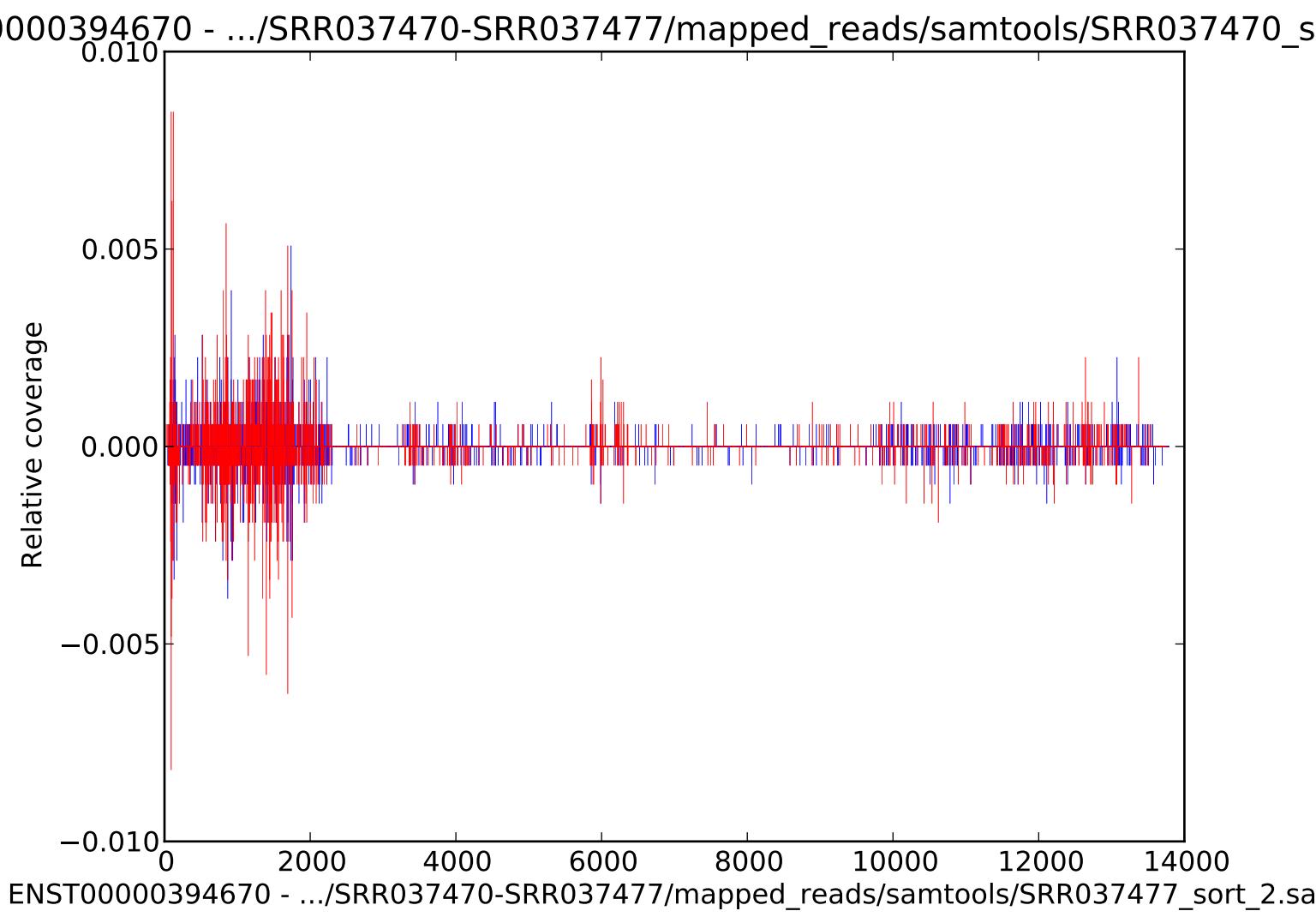
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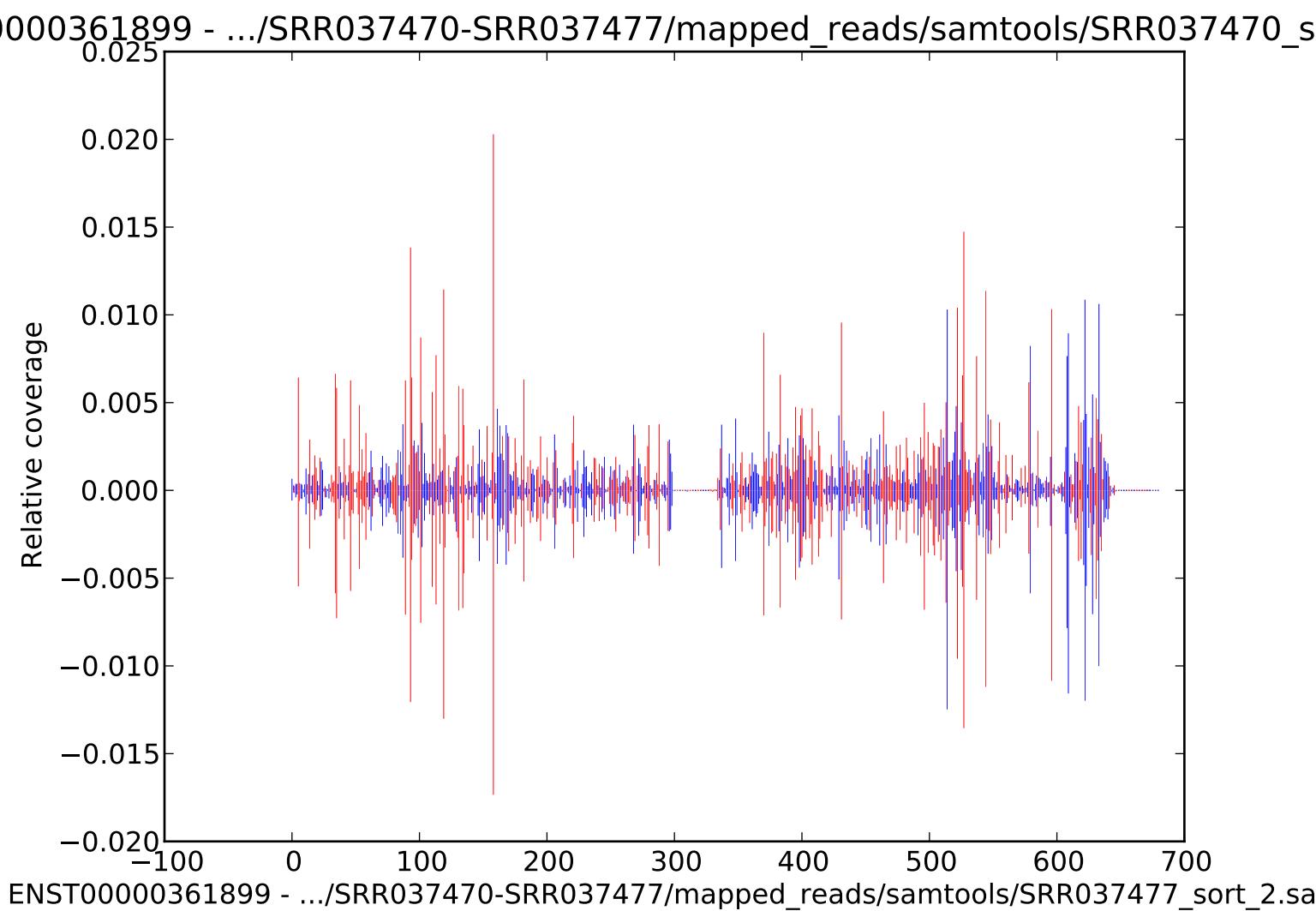
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p=0.000000e+00, Rho=3.578949e-01



p=0.000000e+00, Rho=9.759434e-01



p=0.000000e+00, Rho=3.091841e-01

