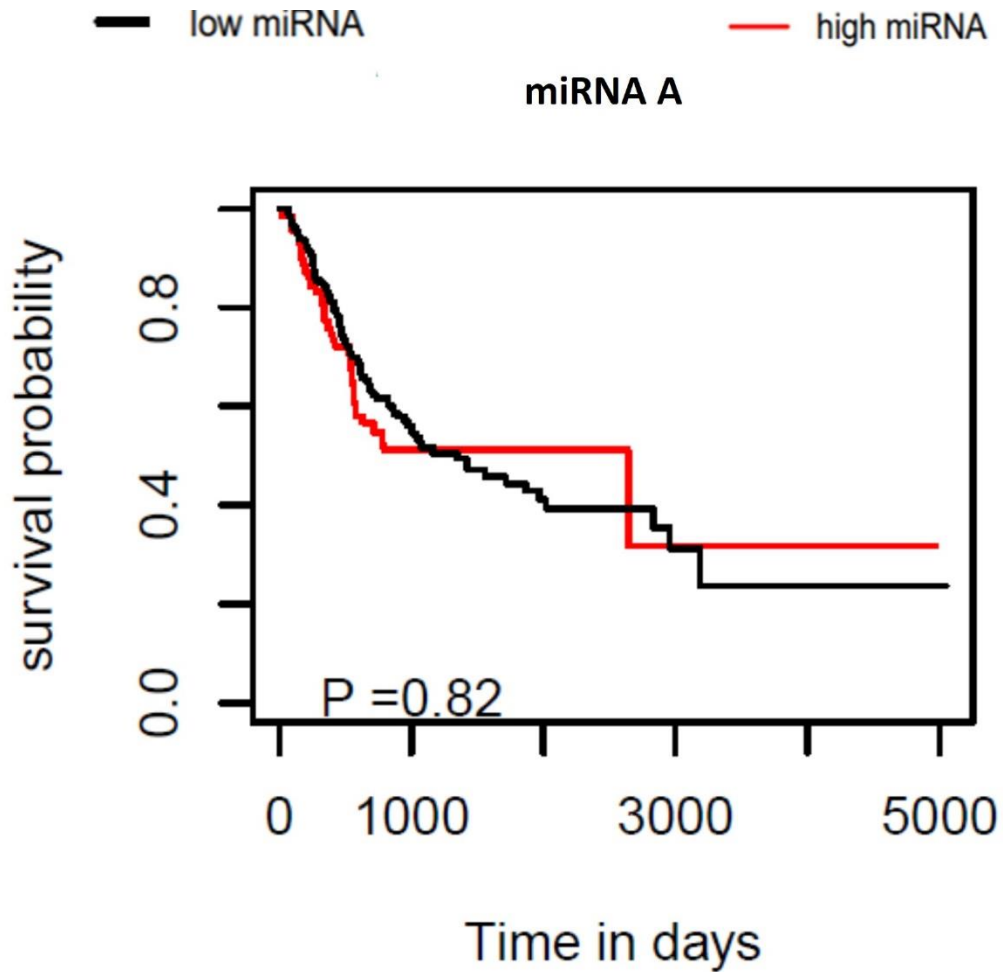


Computing microRNA-gene interaction networks in pan-cancer using miRDriver

Banabithi Bose, Matthew Moravec, and Serdar Bozdag

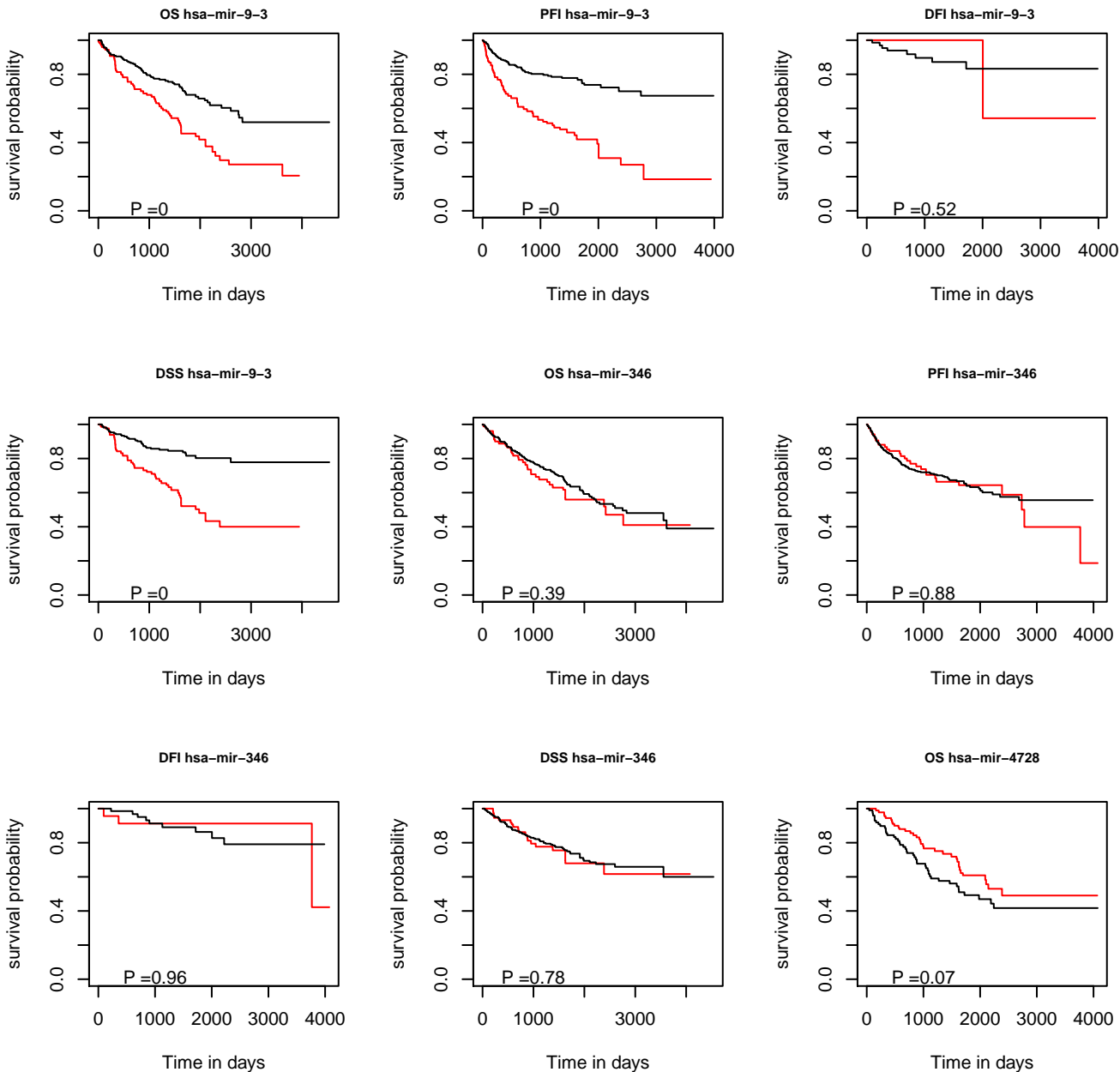
Supplemental Figure S8

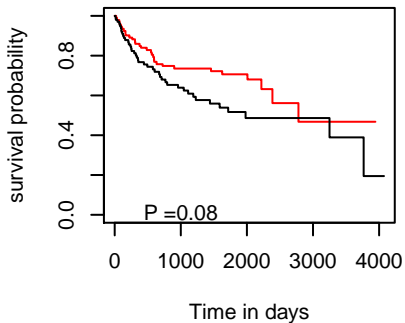
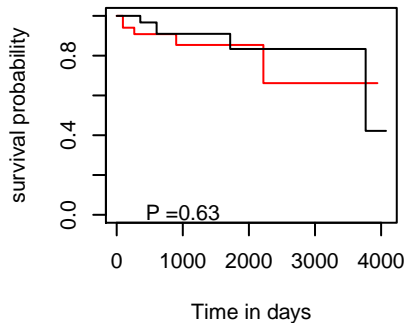
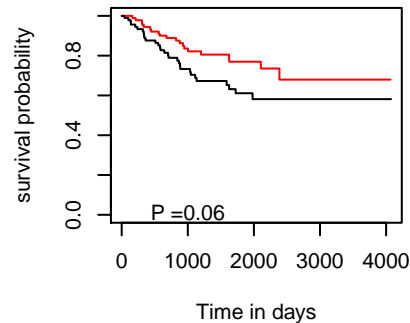
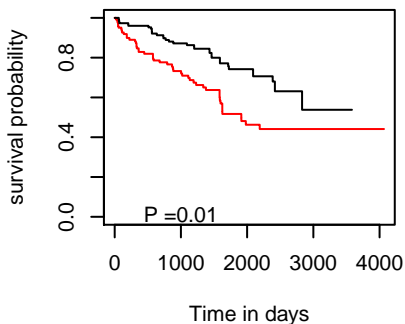
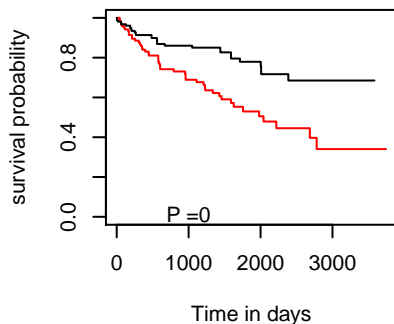
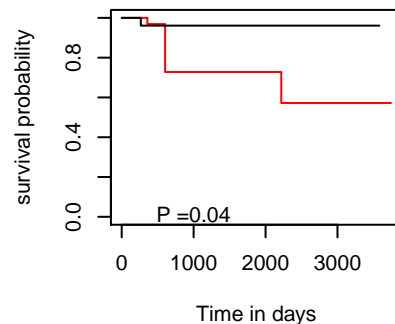
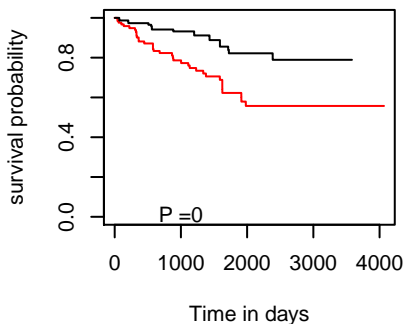
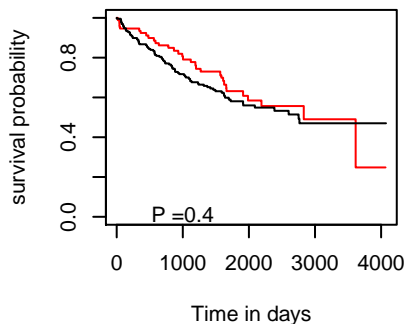
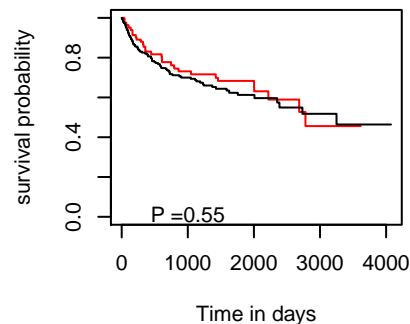


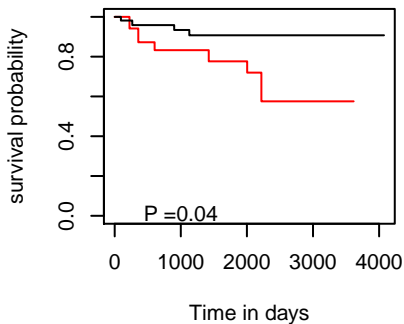
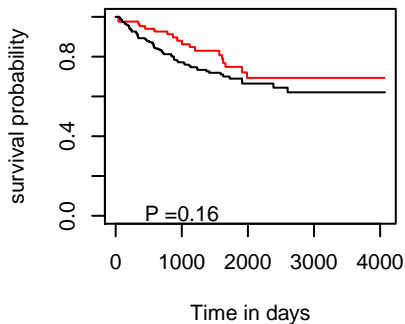
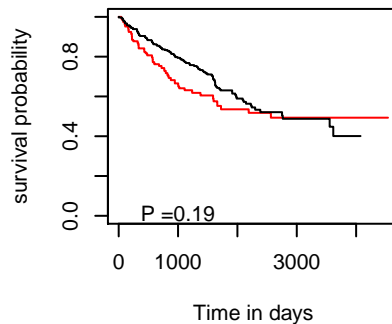
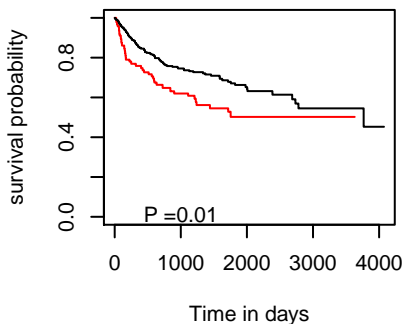
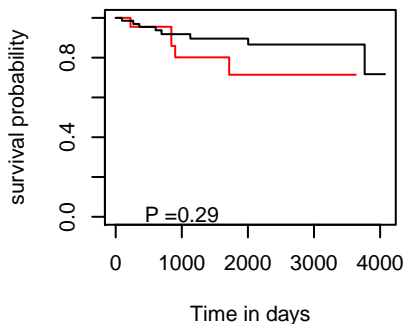
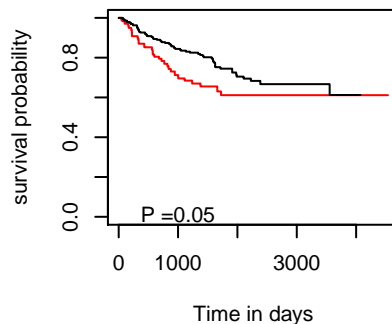
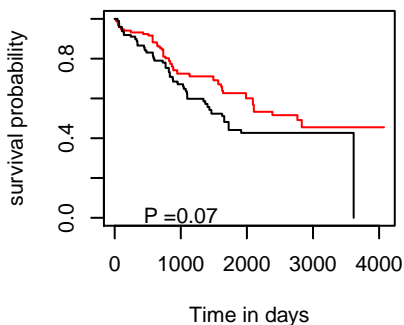
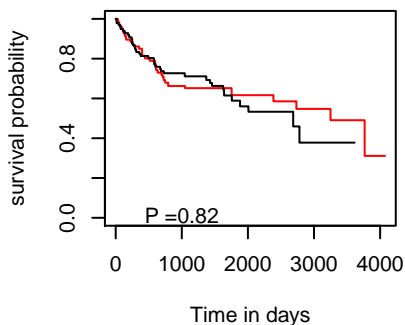
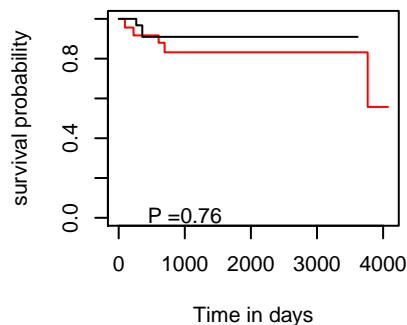
The *Adjusted Kaplan-Meier* survival plots for the computed miRNAs in high and low miRNA expression patient groups.

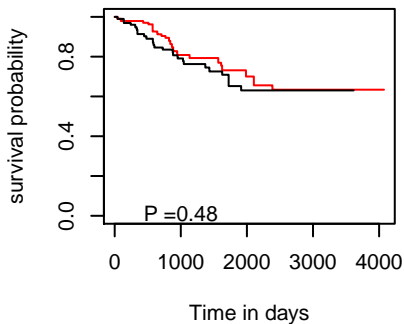
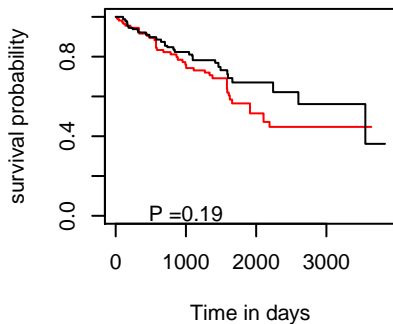
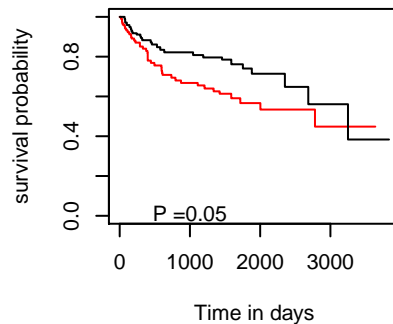
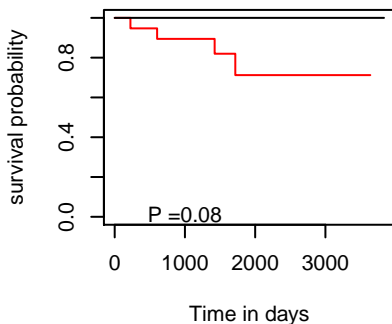
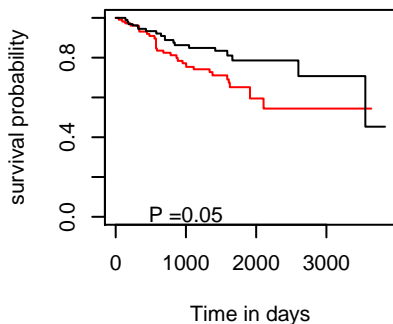
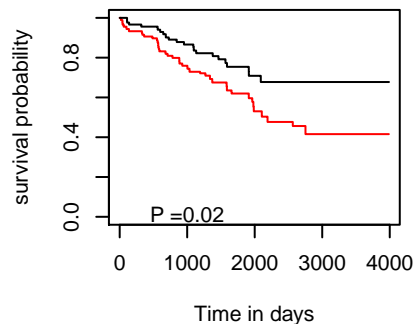
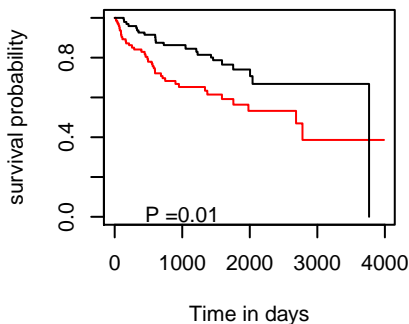
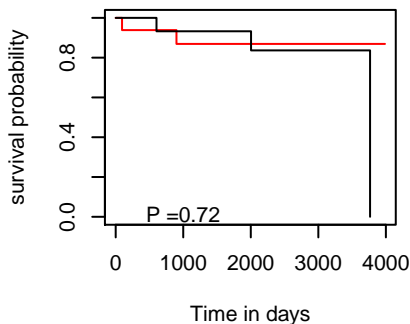
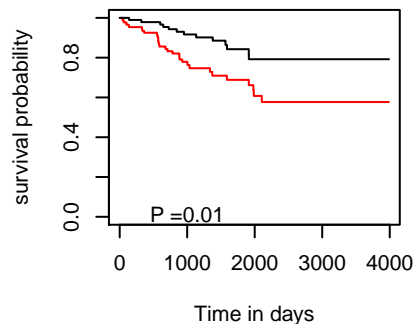
Supplemental Figure S8

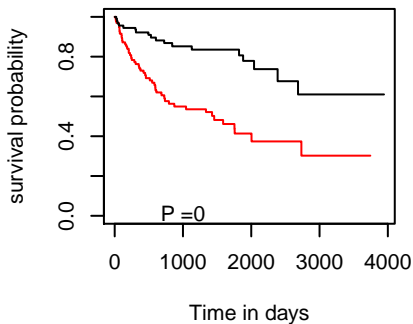
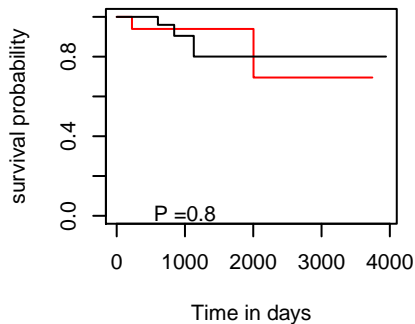
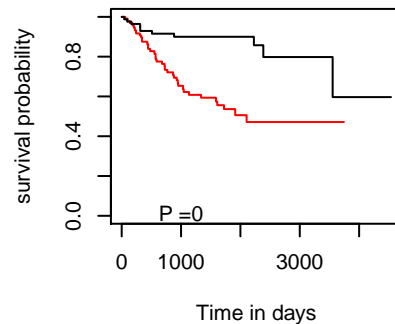
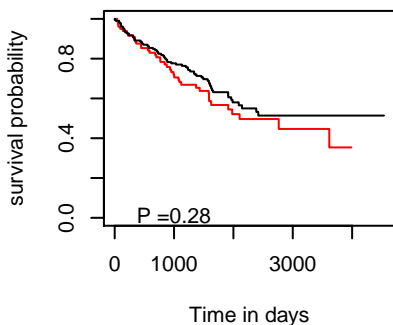
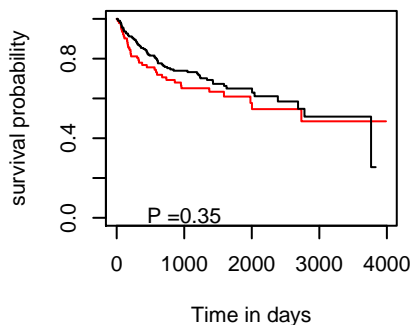
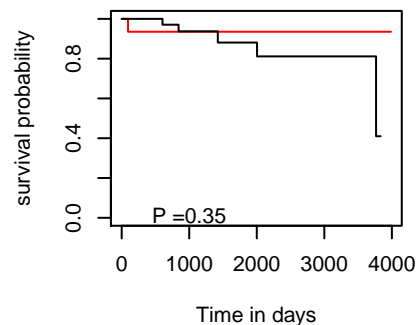
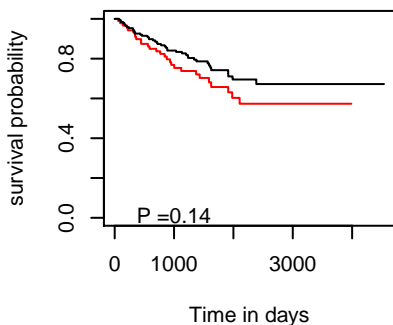
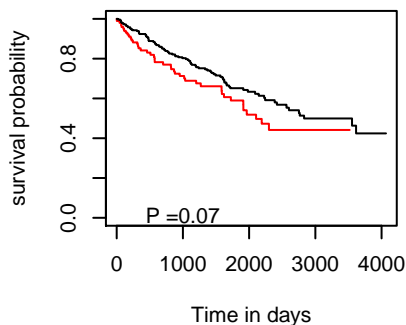
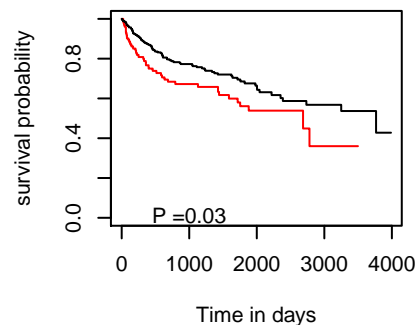
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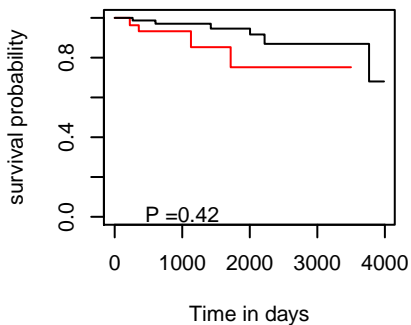
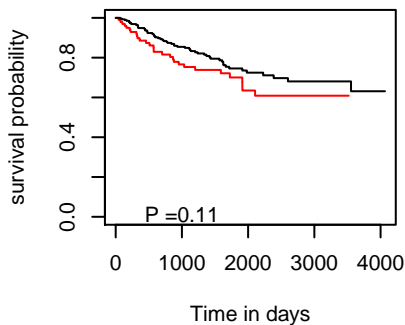
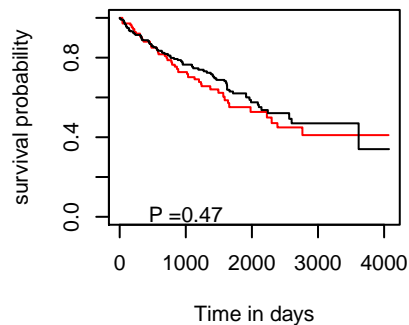
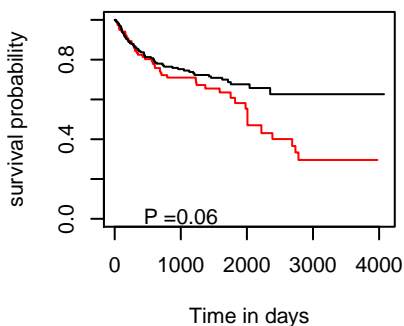
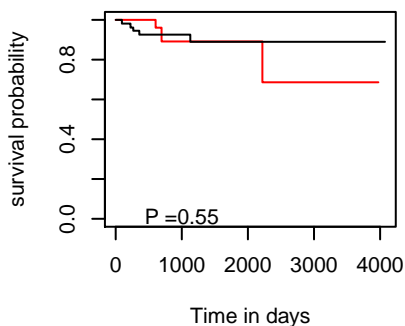
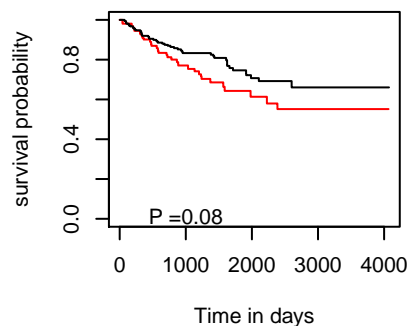
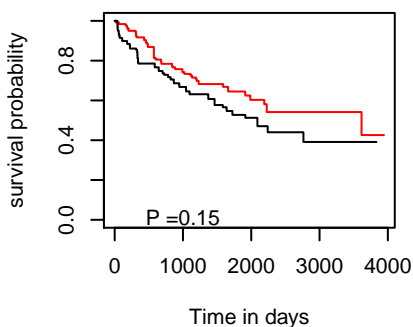
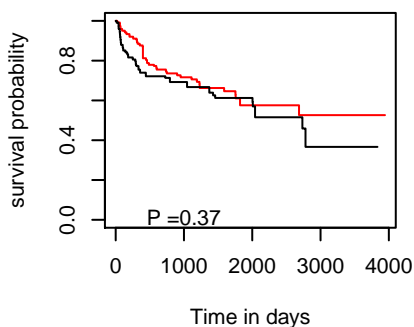
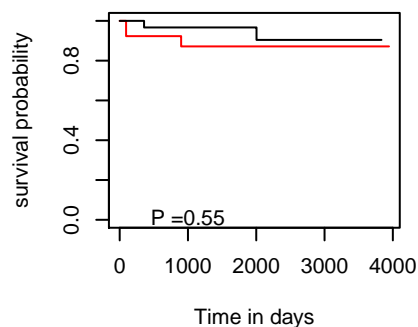


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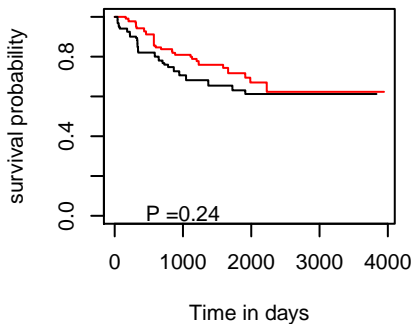
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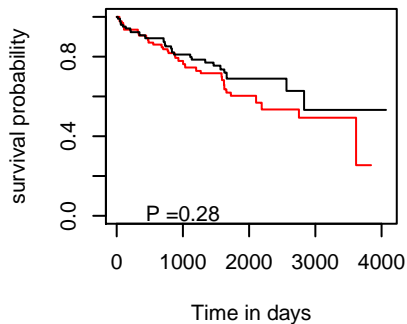
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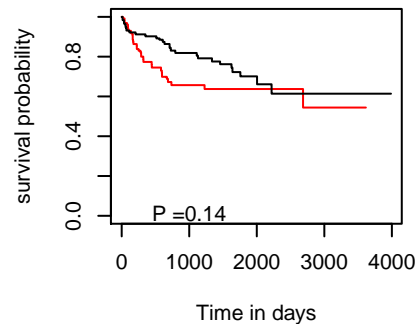
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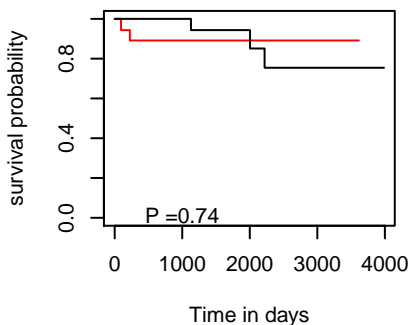
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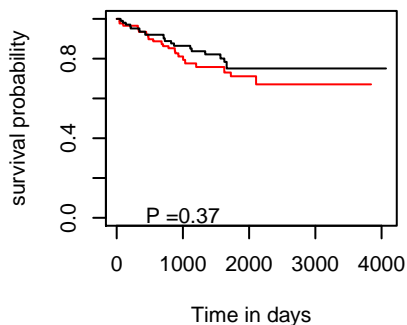
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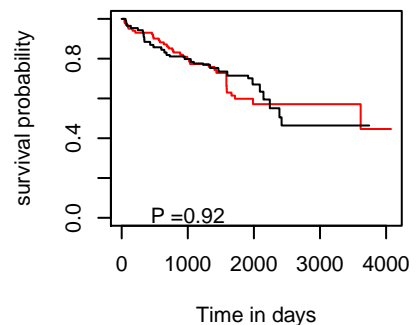
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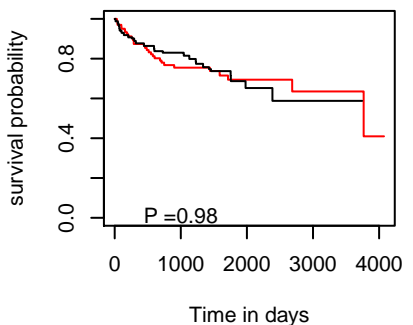
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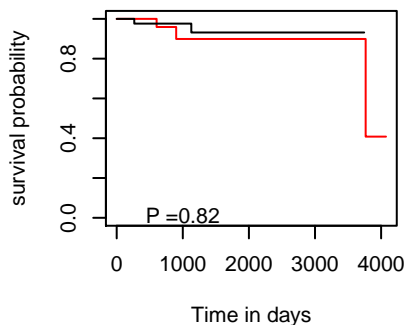
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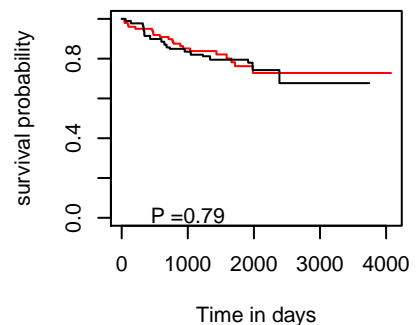
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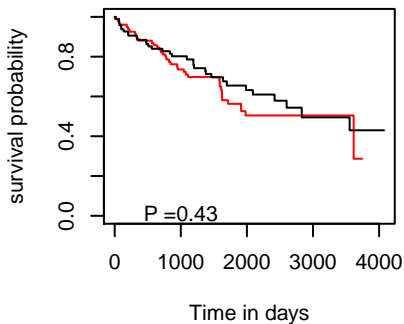
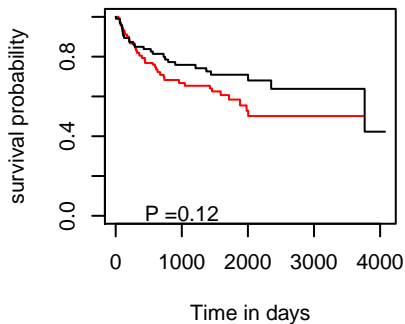
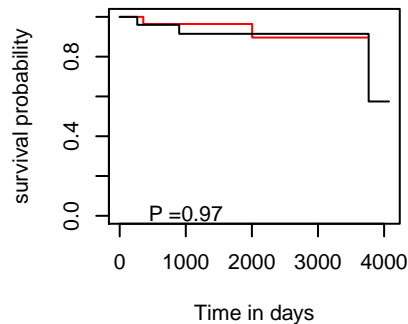
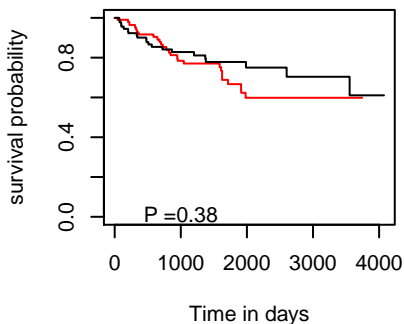
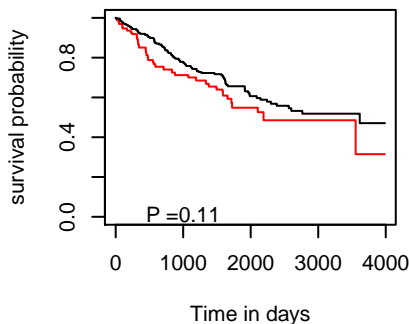
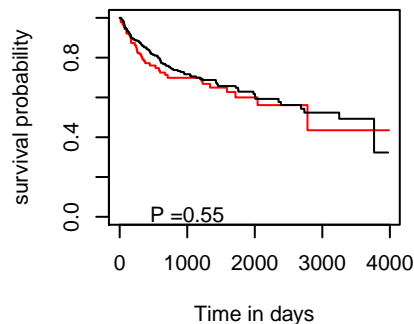
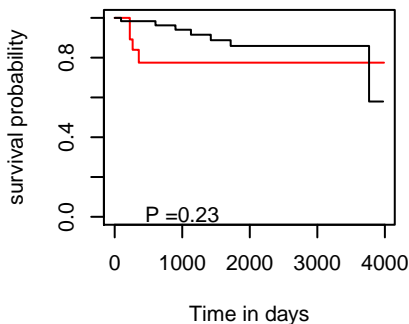
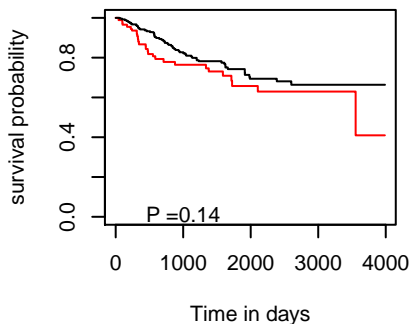
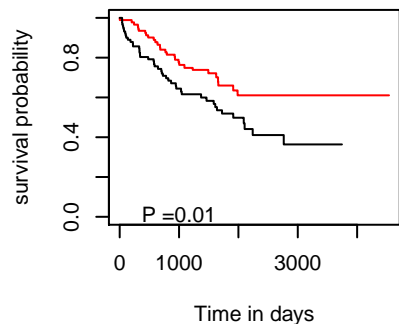


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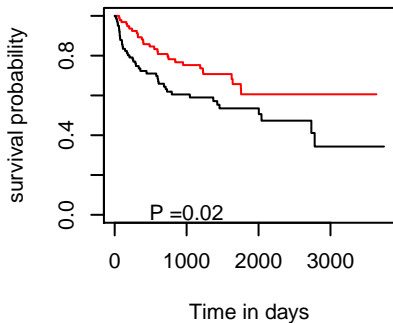


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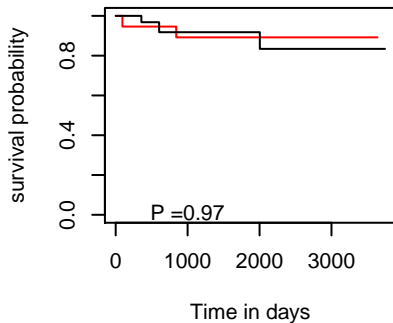


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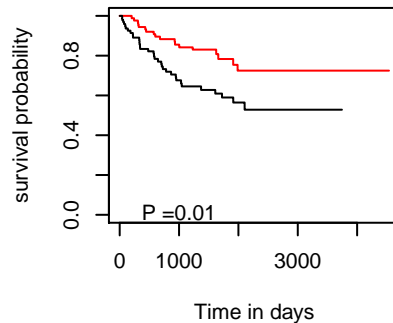
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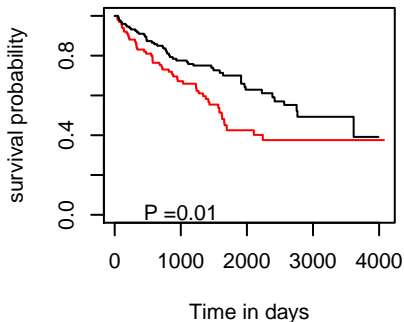
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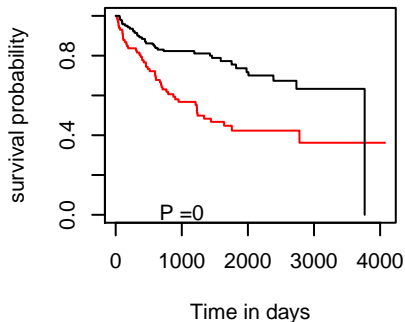
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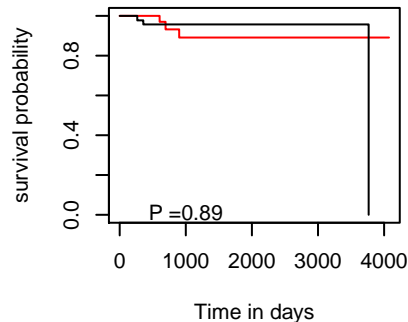
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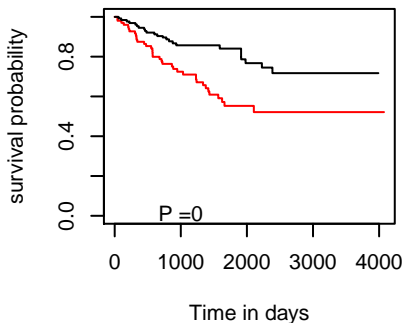
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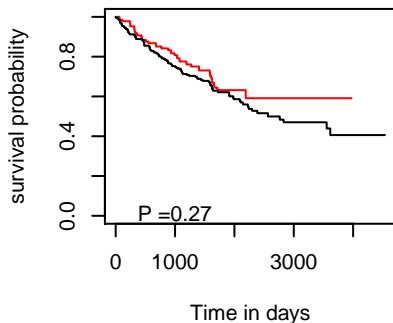
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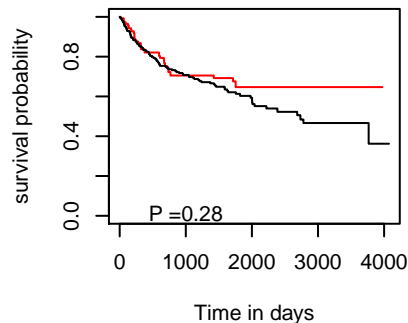
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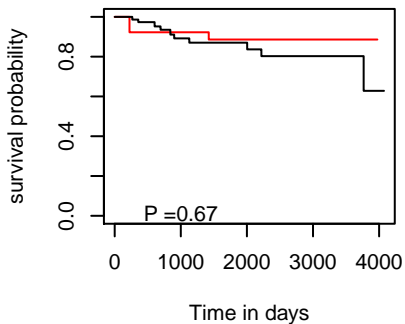
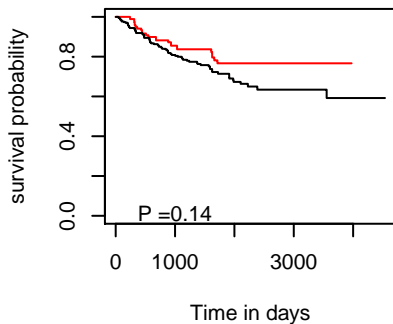
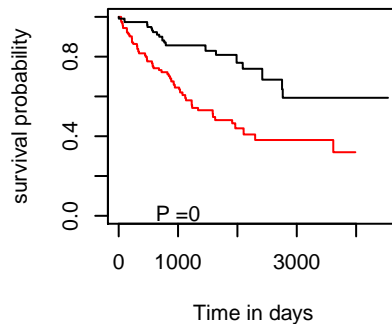
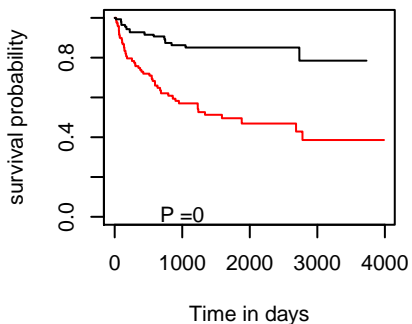
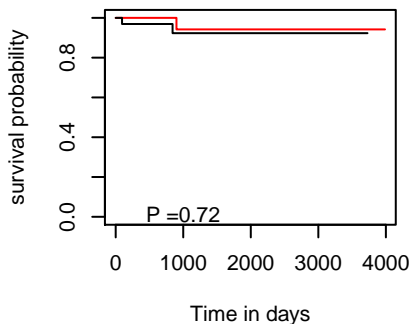
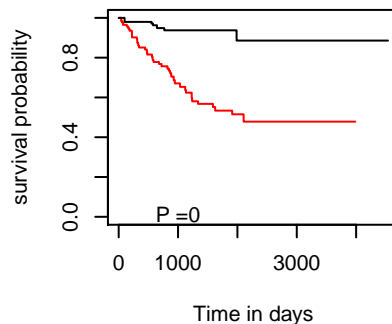
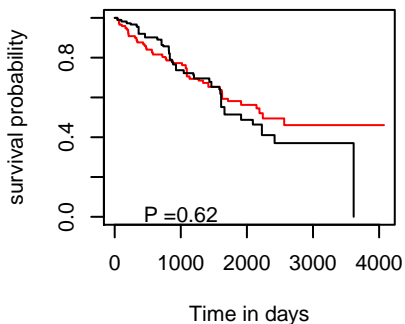
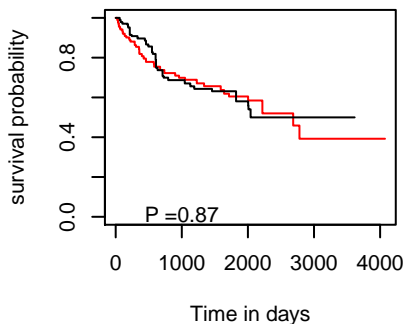
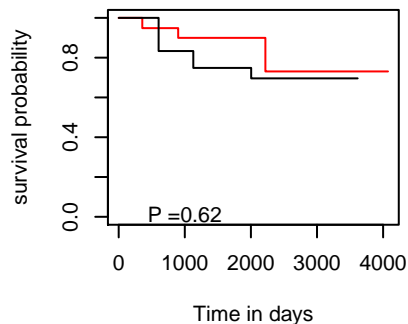


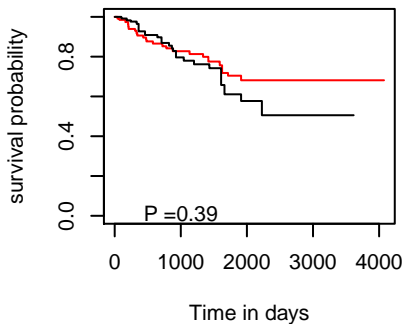
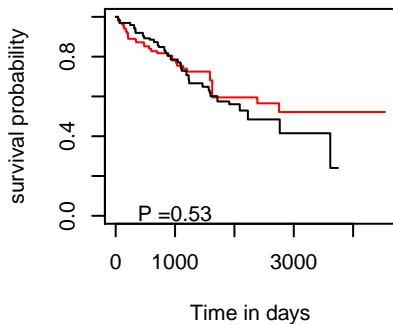
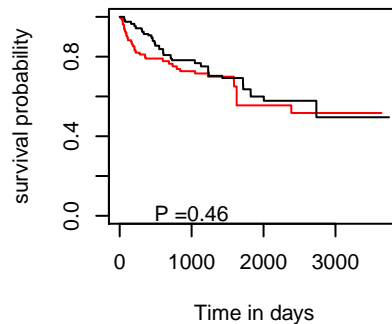
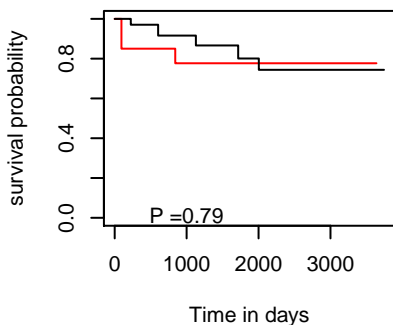
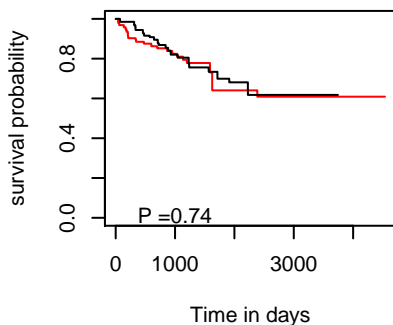
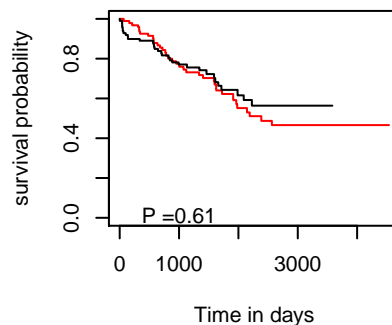
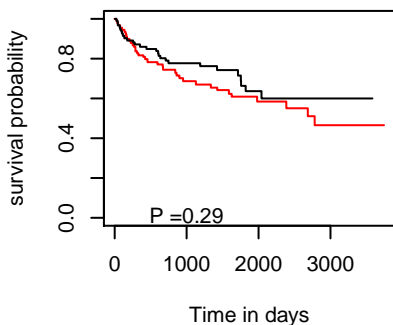
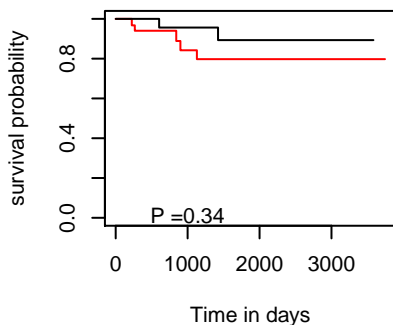
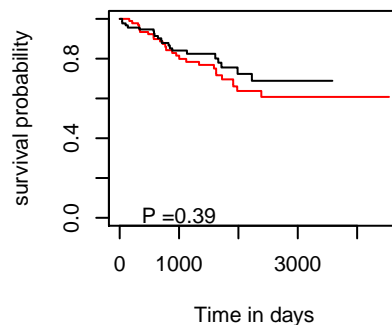
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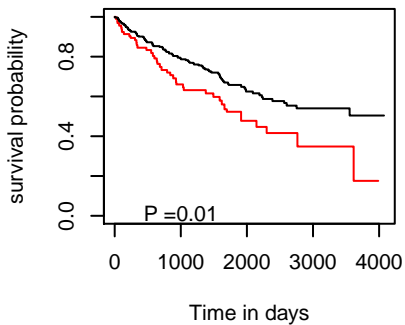
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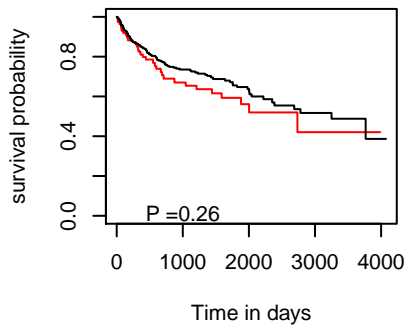
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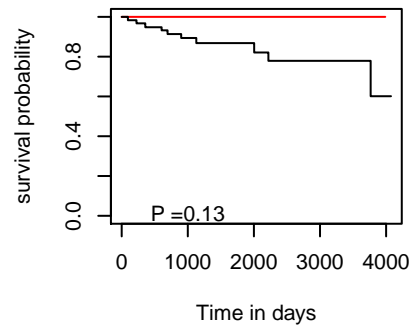
OS hsa-mir-4777



PFI hsa-mir-4777



DFI hsa-mir-4777



DSS hsa-mir-4777

