

Figure S1. Protein interaction analysis by STRING database (left) and KEGG pathway clustering analysis (right) in cell lysis group candidate proteins.

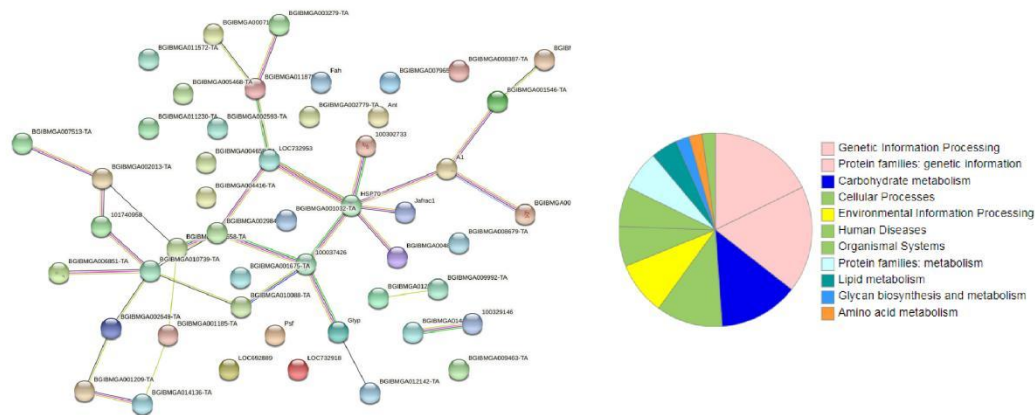


Figure S2. Protein interaction analysis by STRING database (left) and KEGG pathway clustering analysis (right) in live cell group candidate proteins.

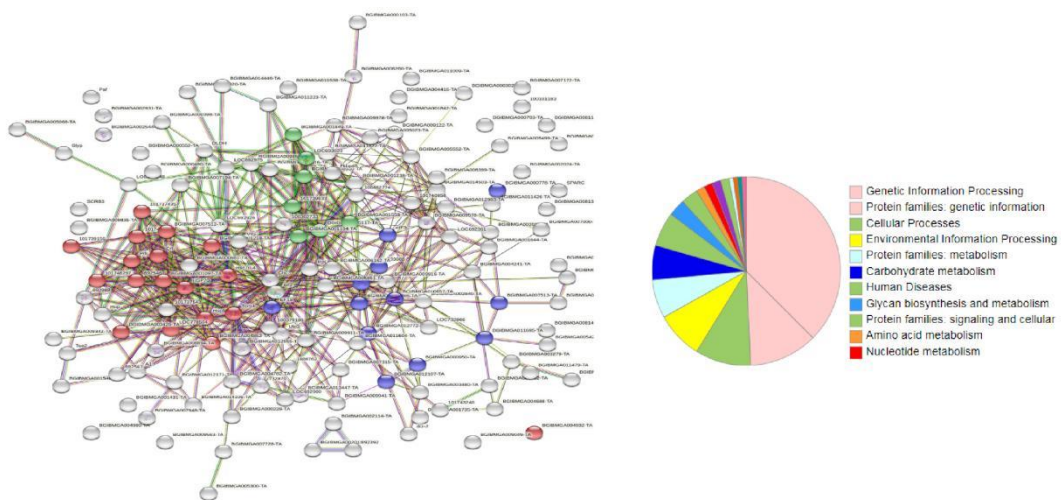
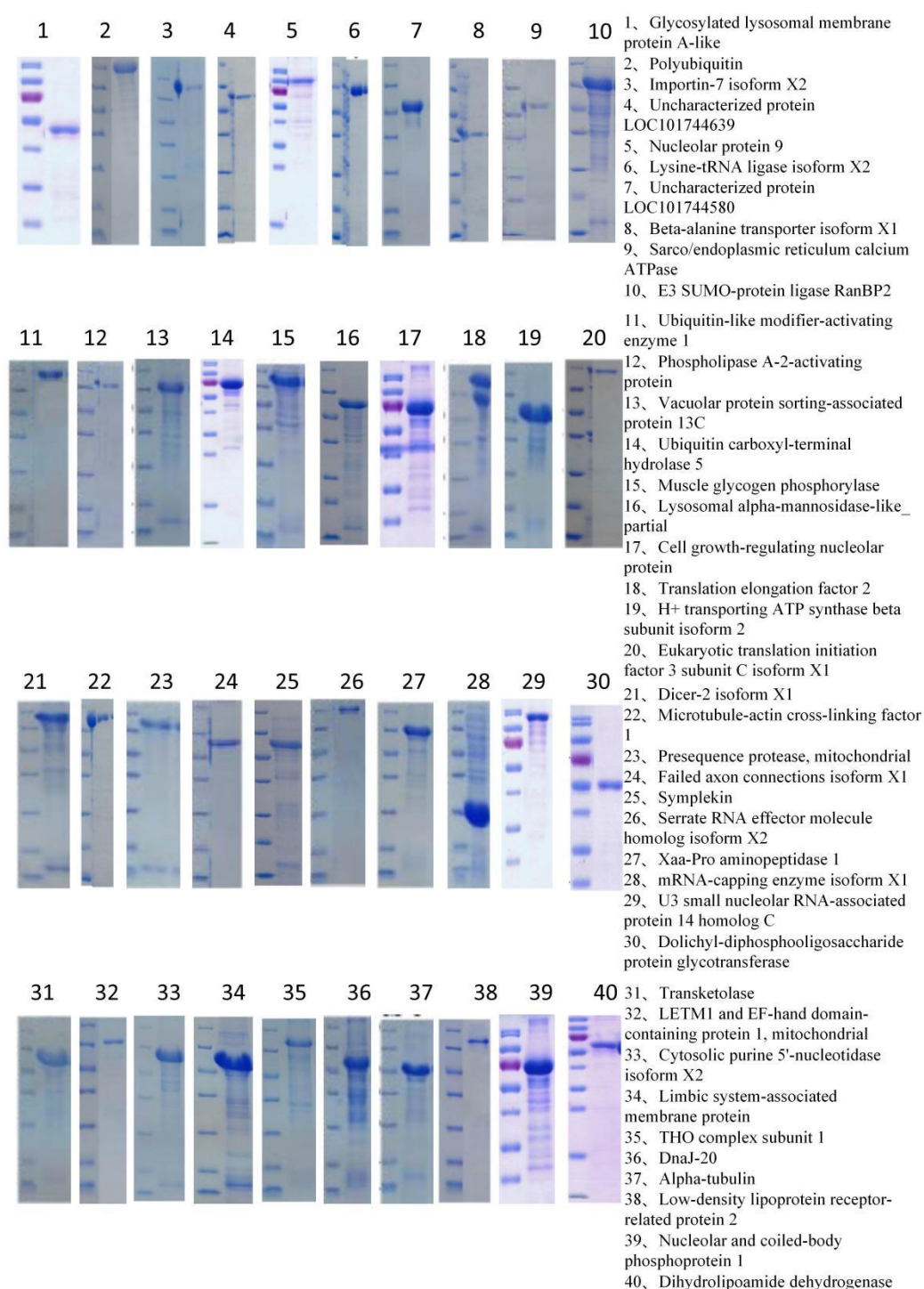


Figure S3. Expression and purification of candidate protein.



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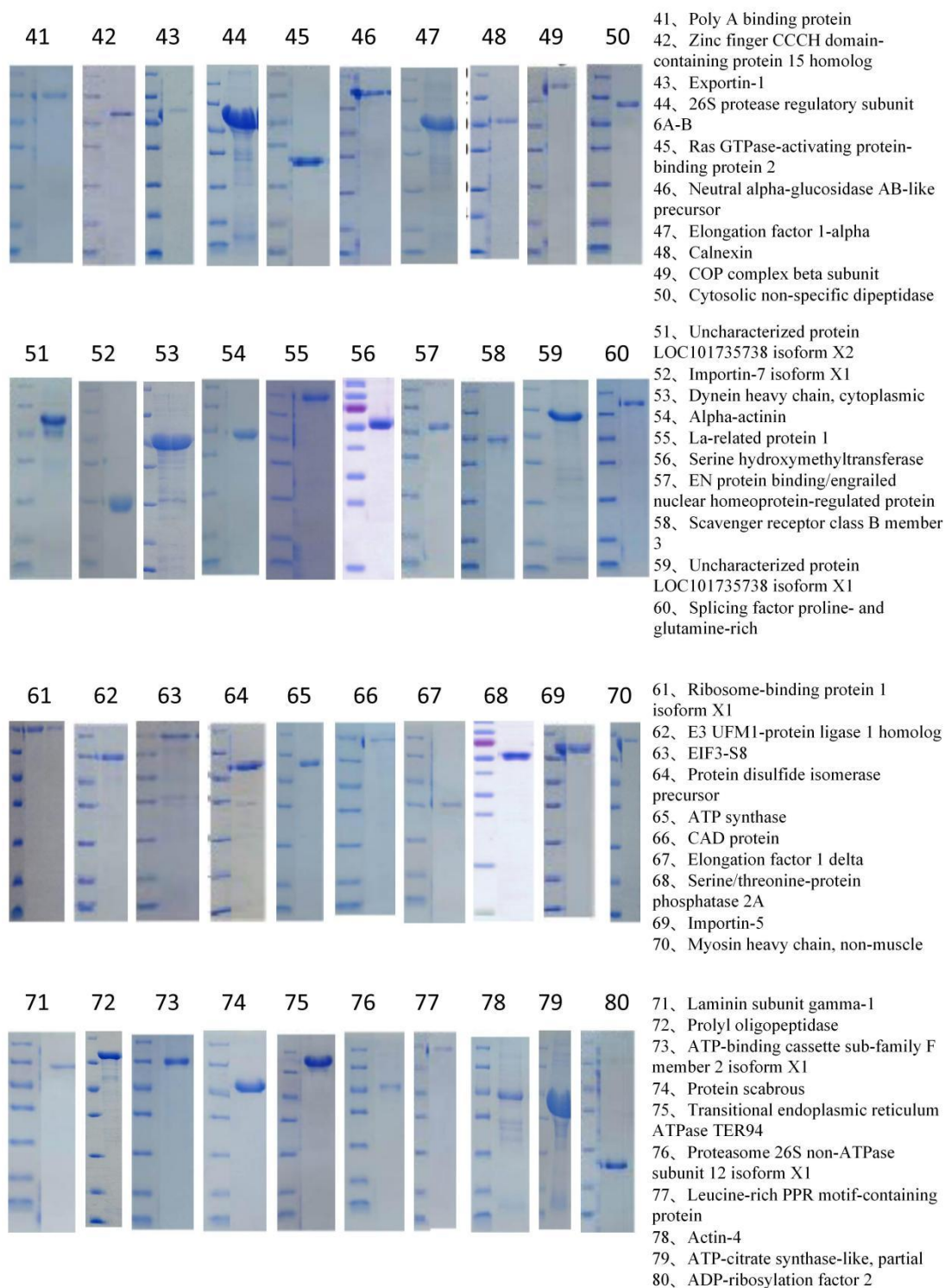
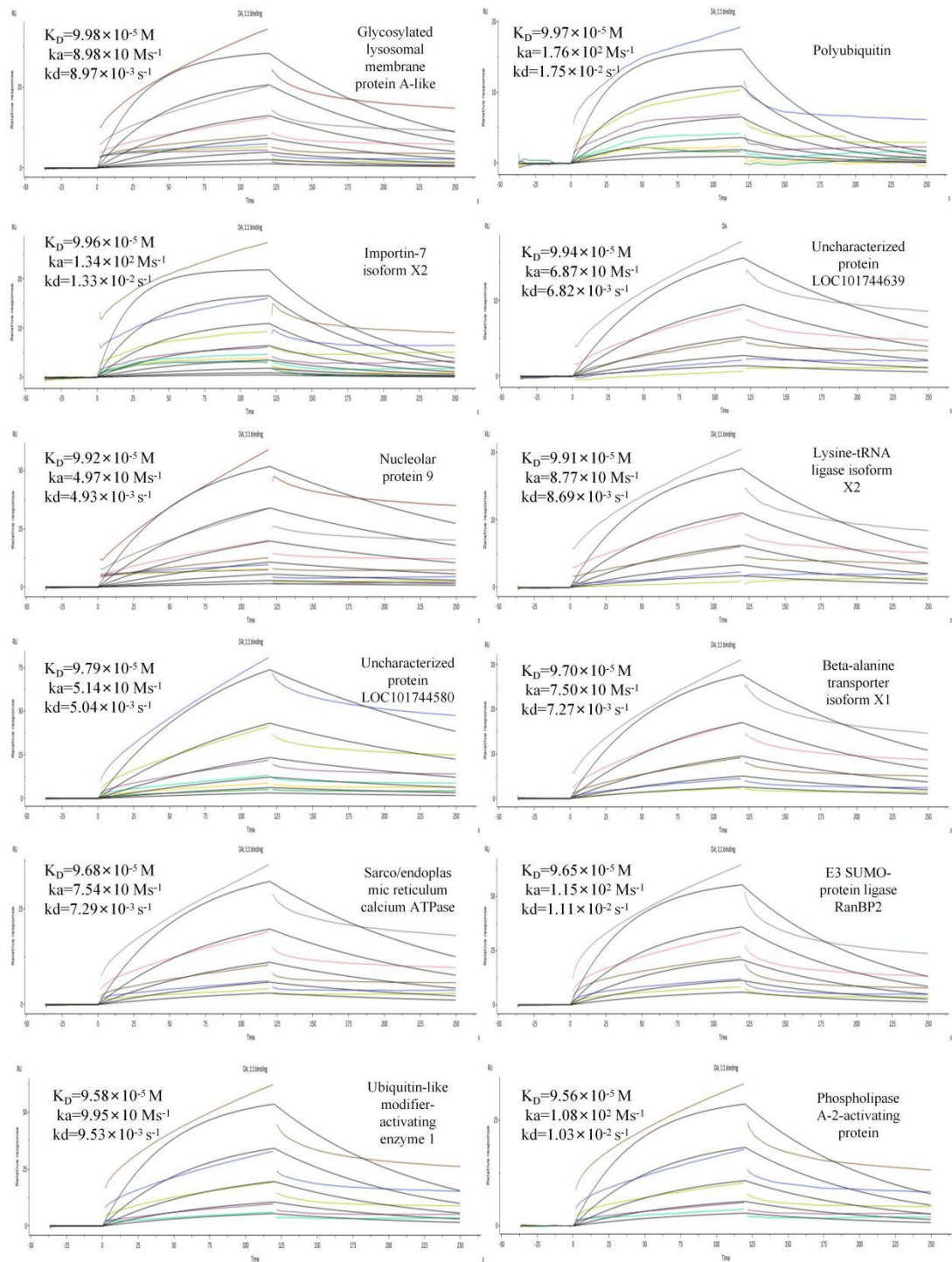
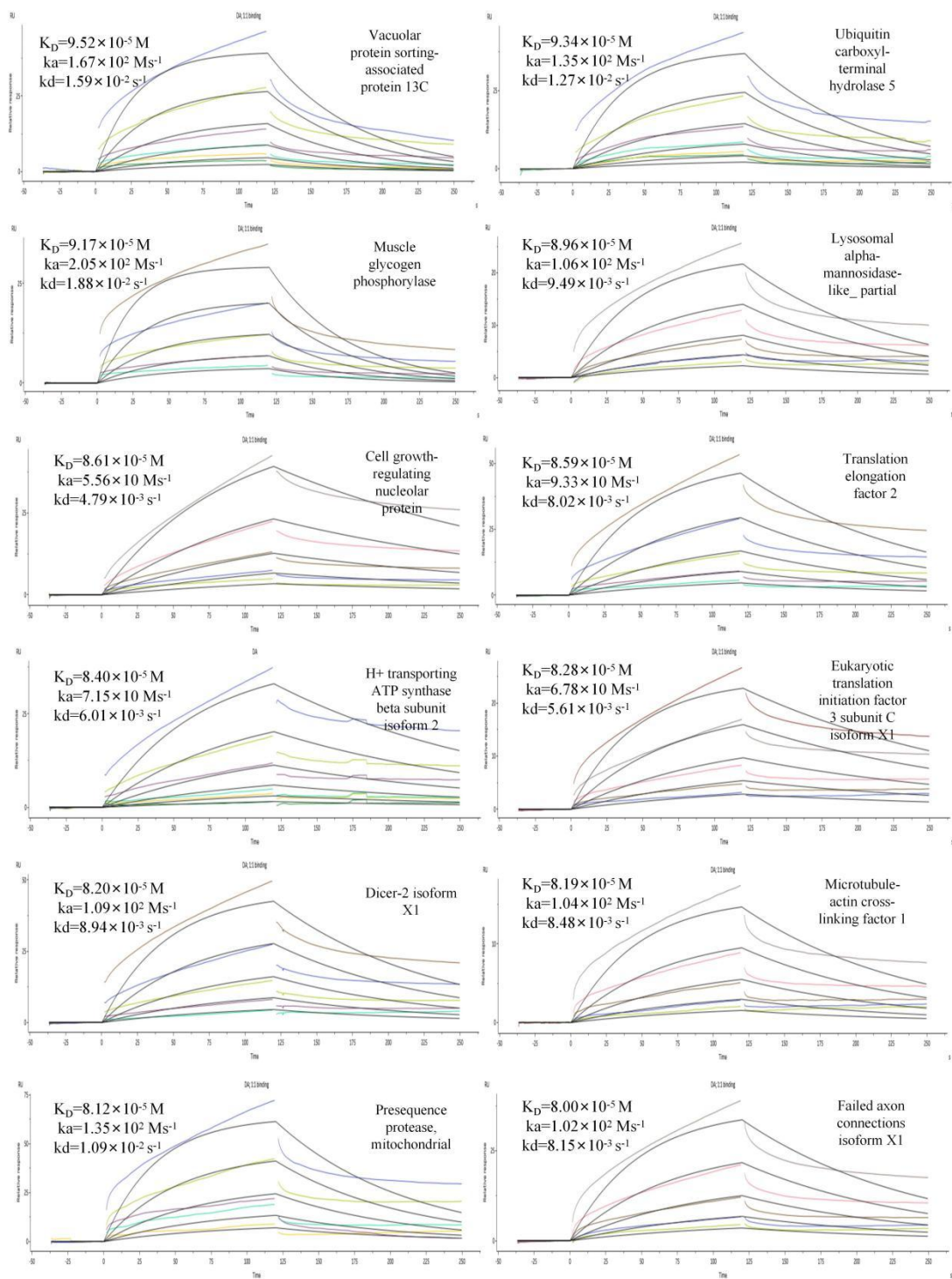


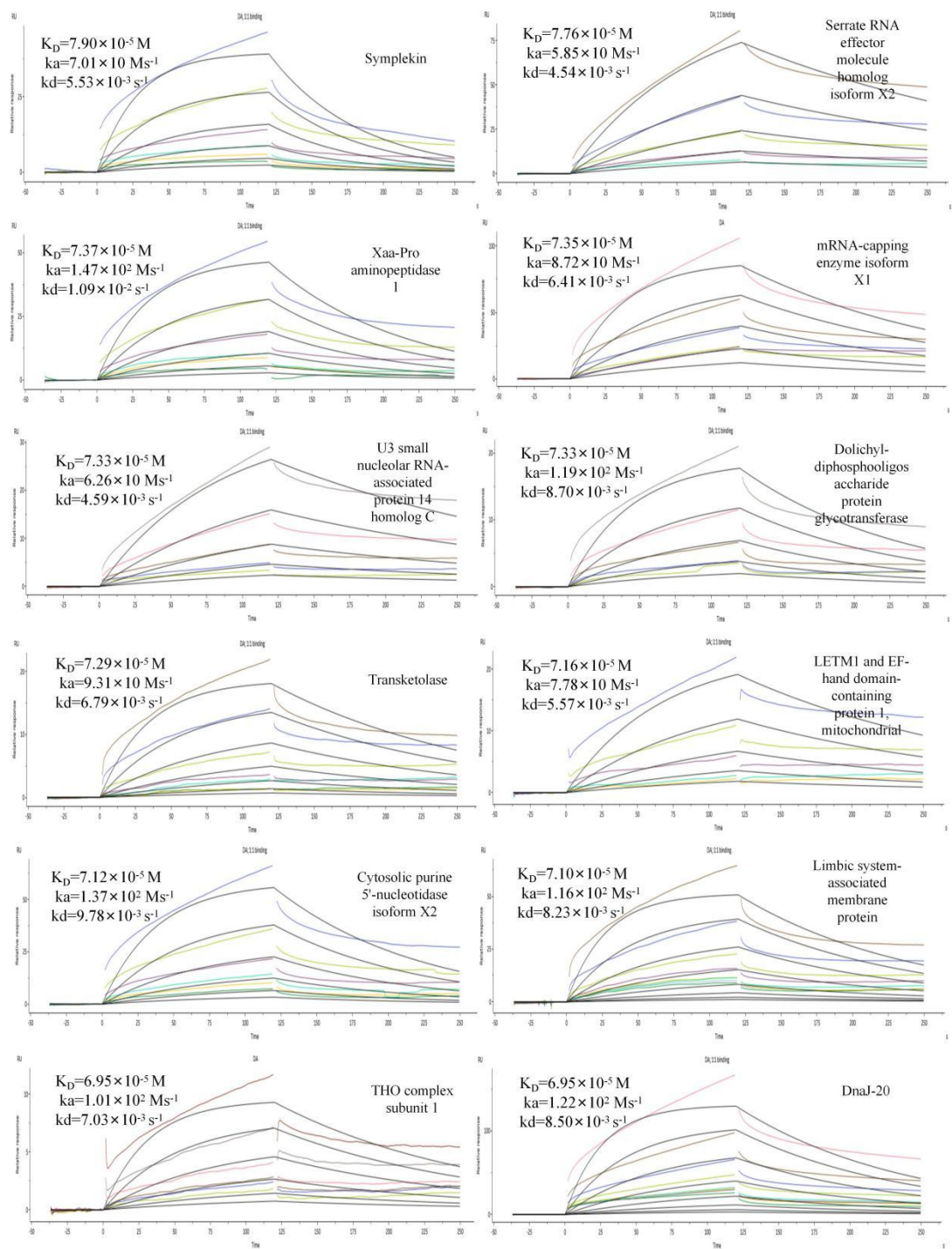
Figure S4. Detailed SPR results of interaction of DA with candidate proteins.



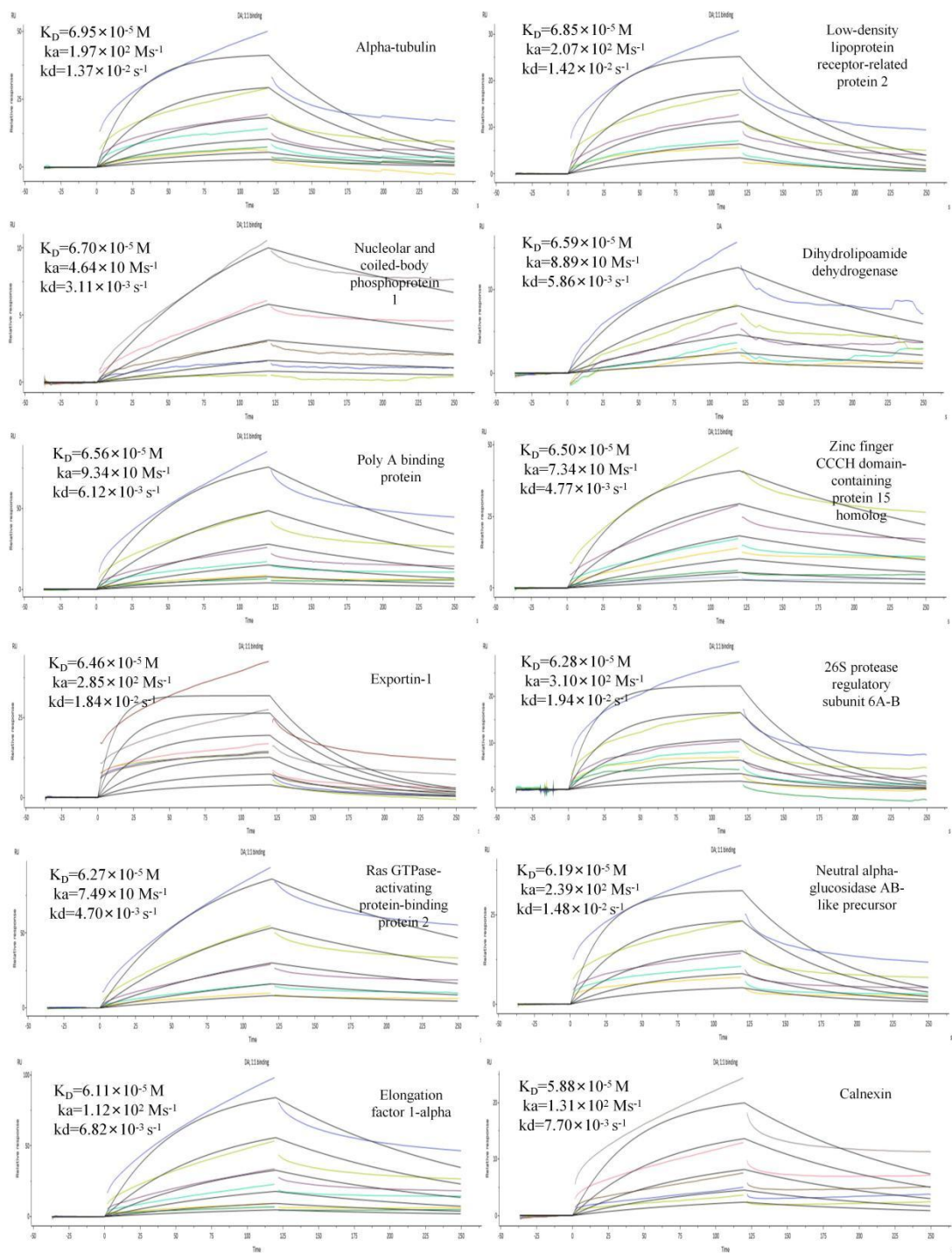
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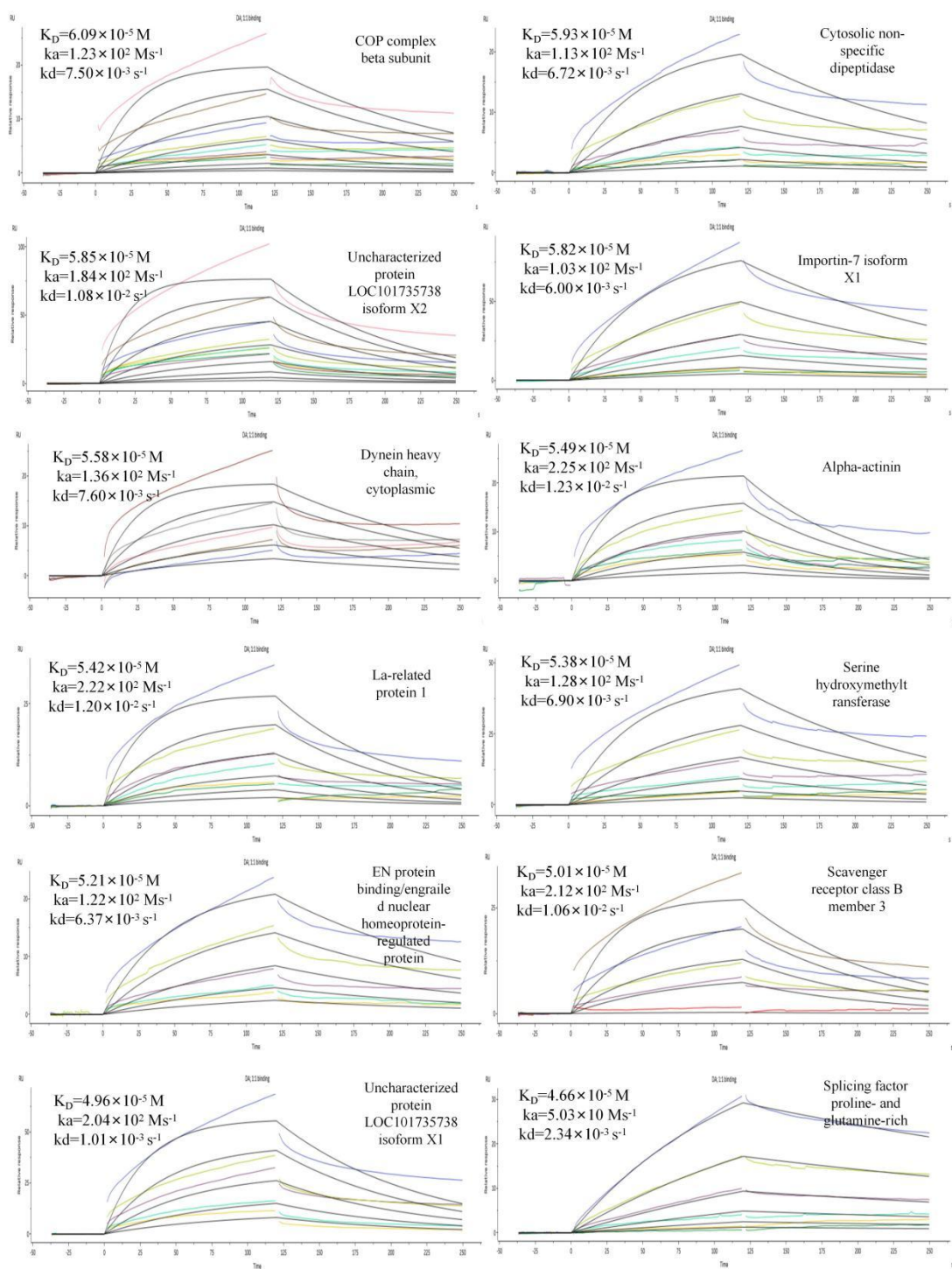
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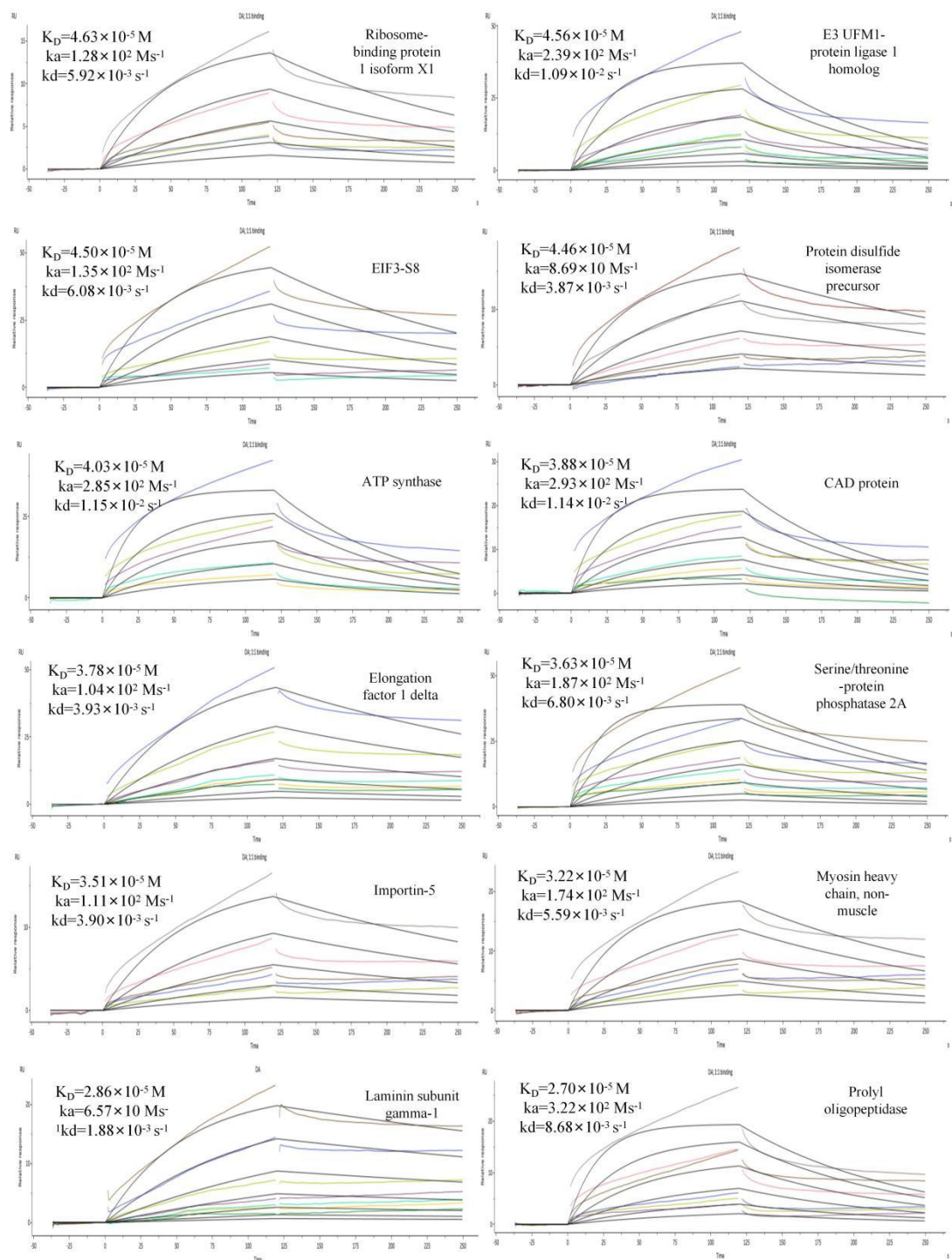
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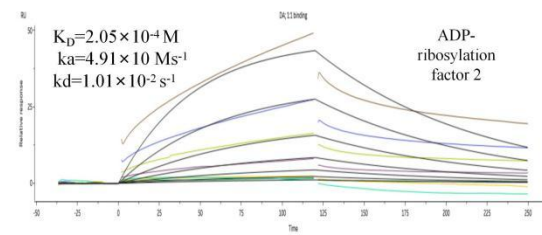
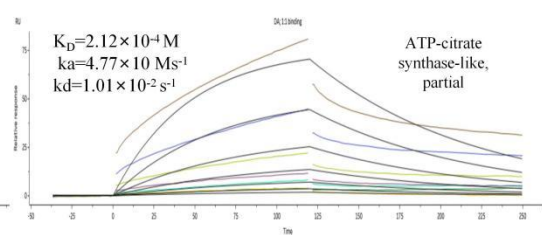
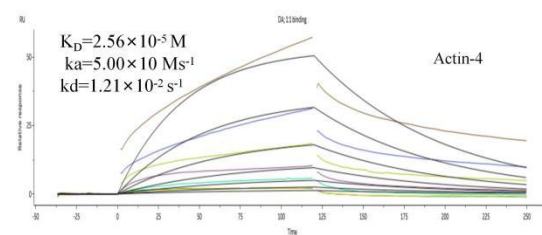
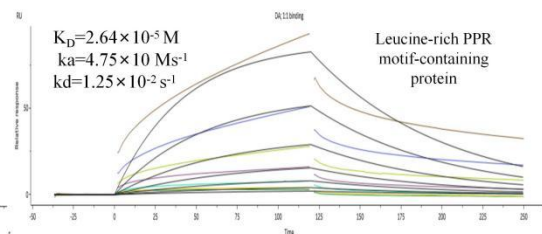
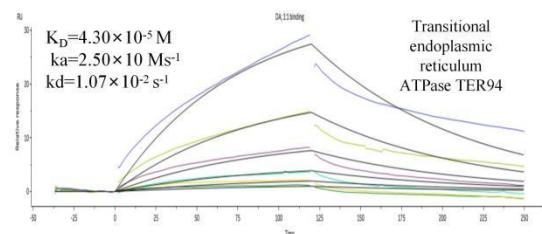
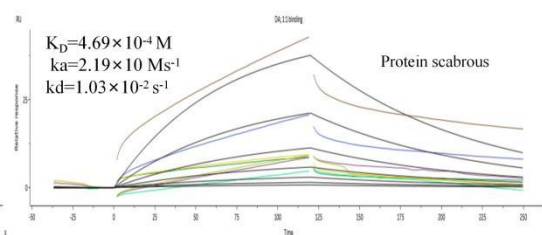
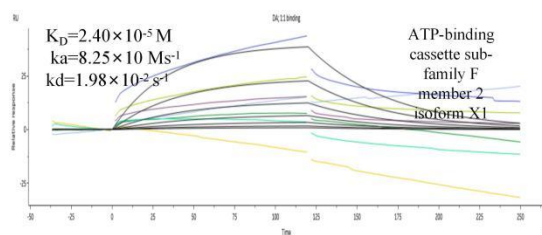
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Original images

Figure 1A

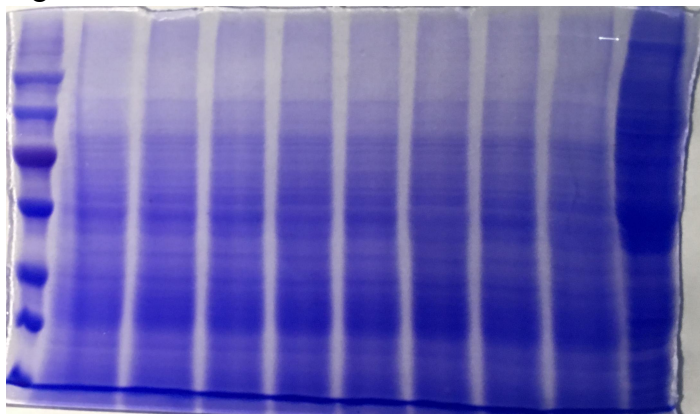
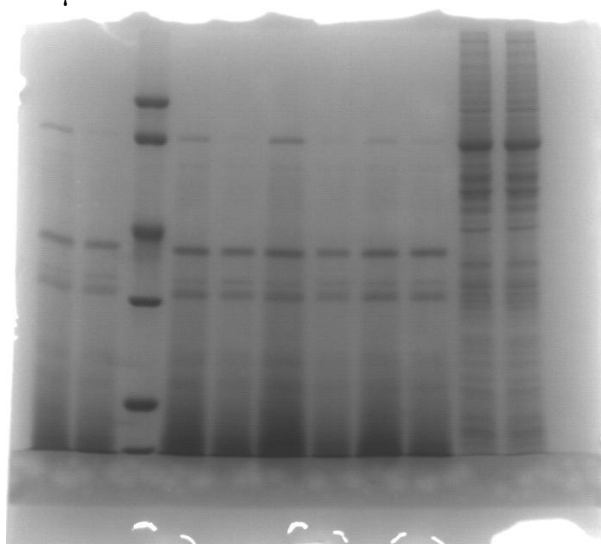
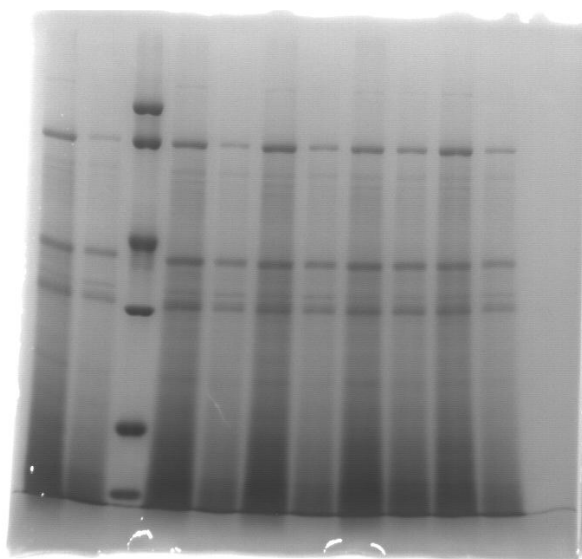


Figure 1B

200 μ M



20 μ M



2 μ M

