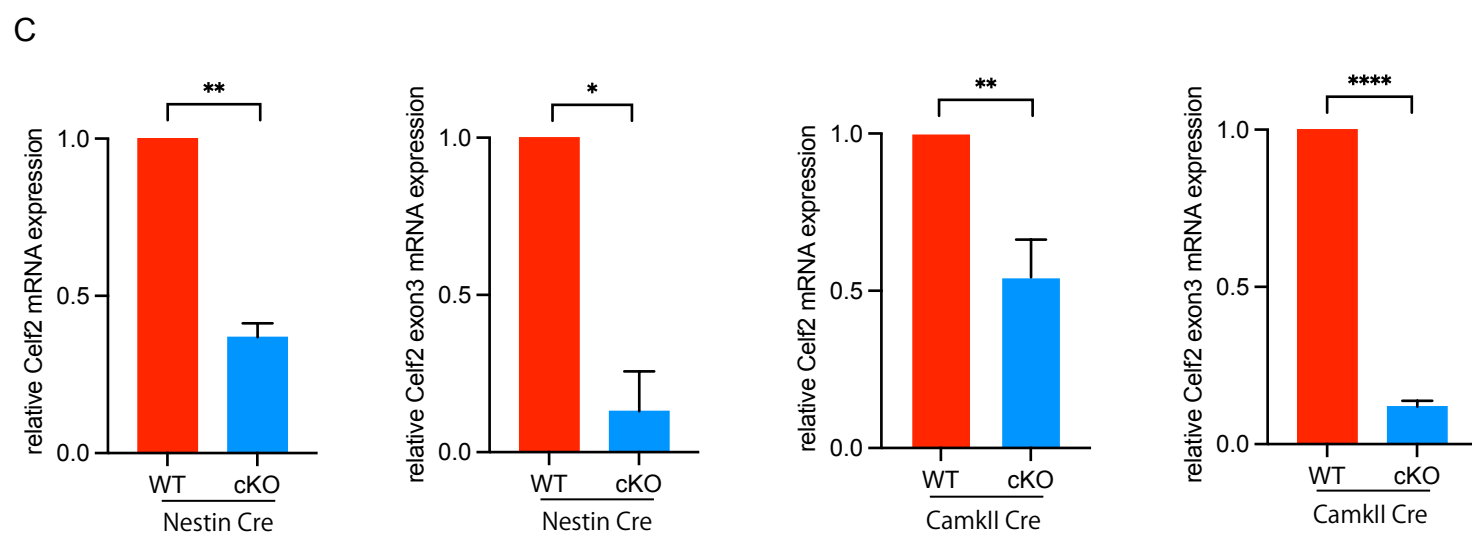
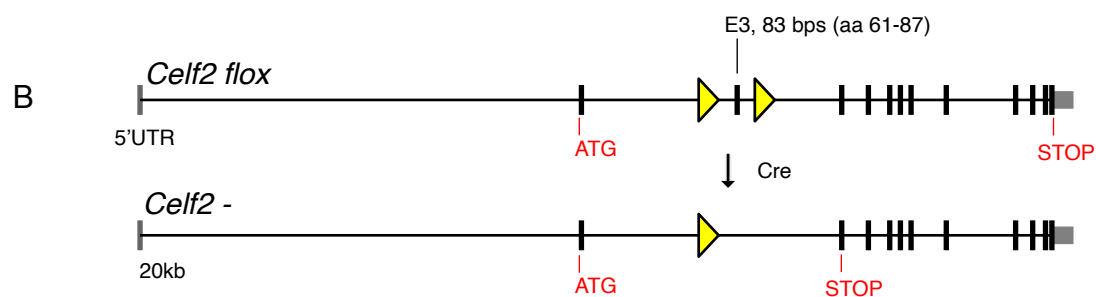
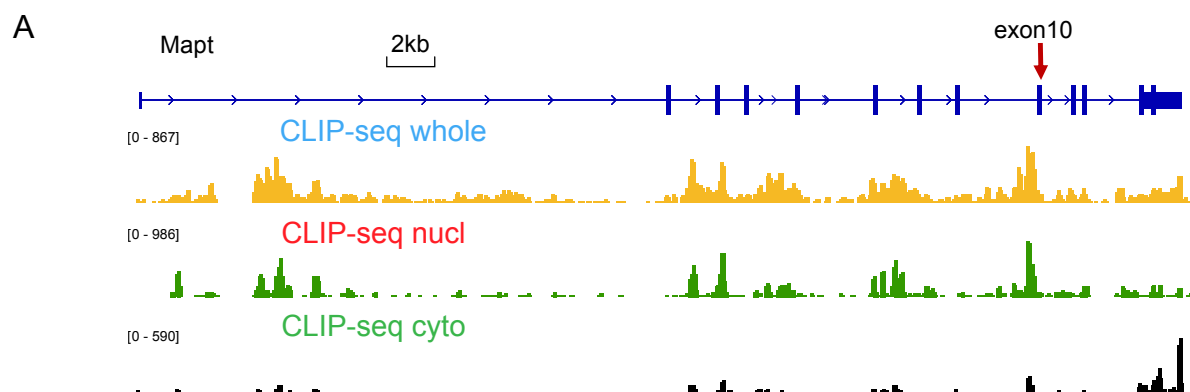
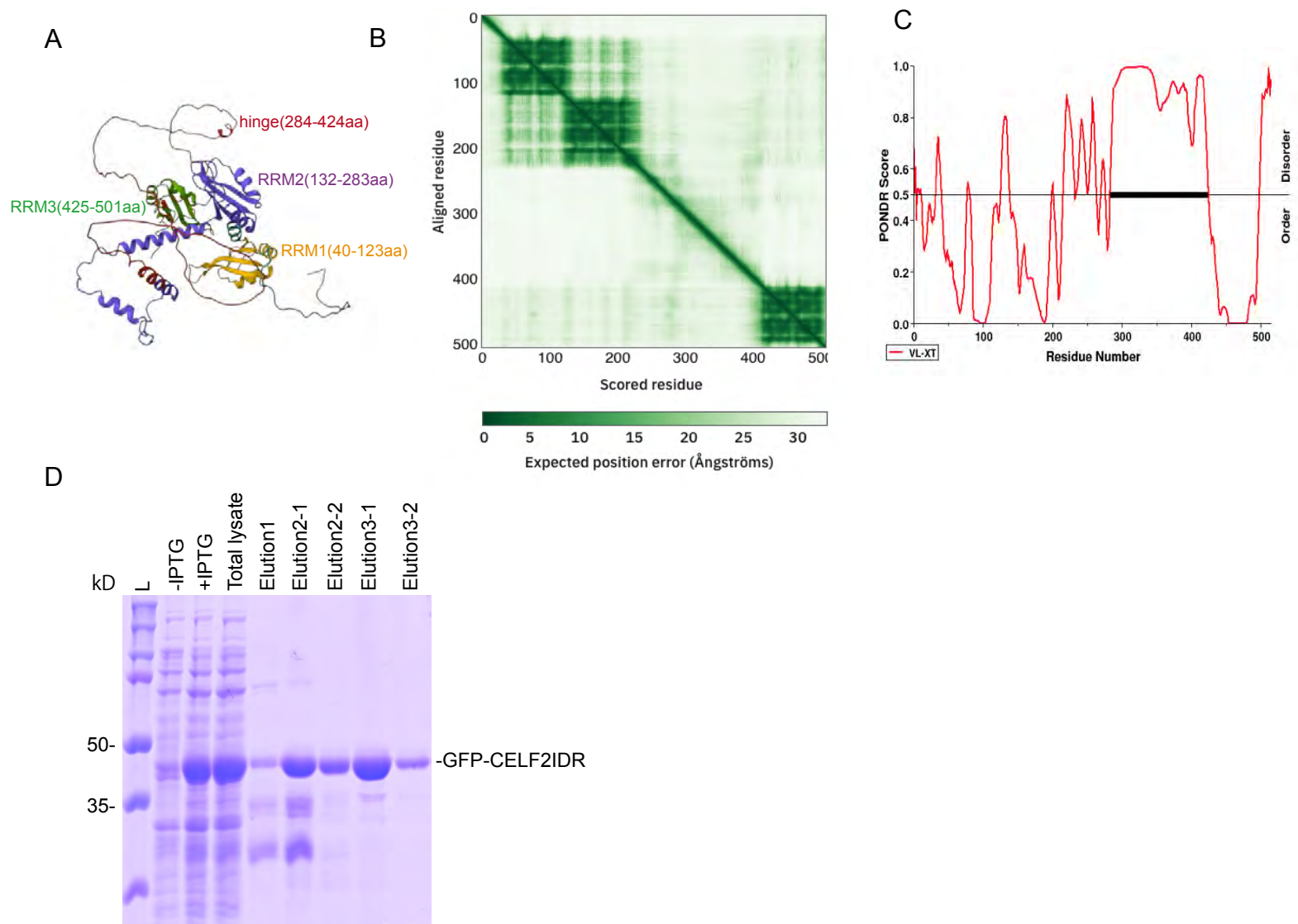
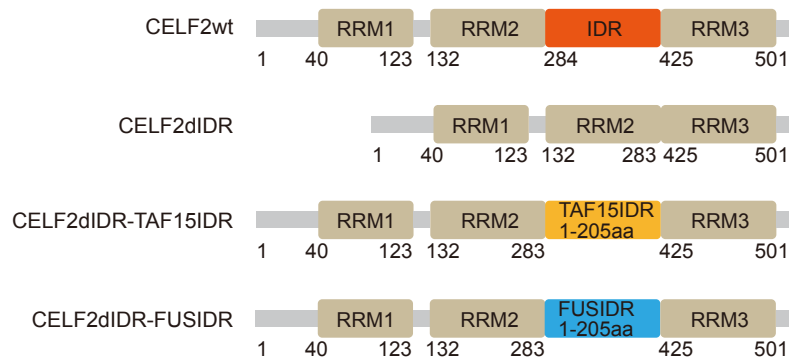


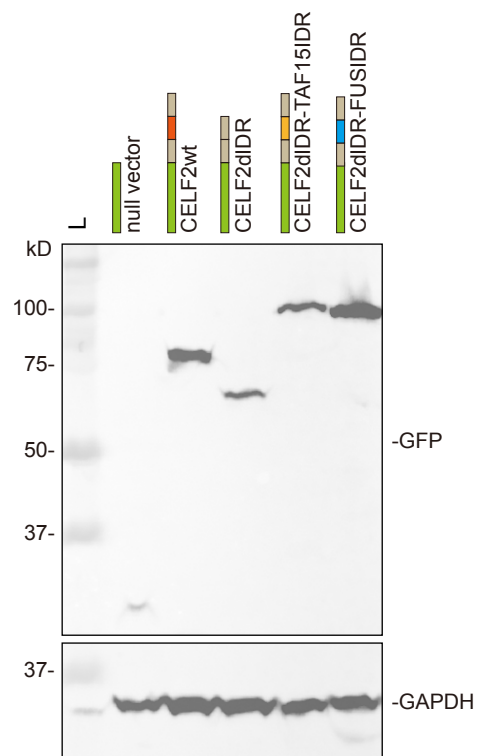




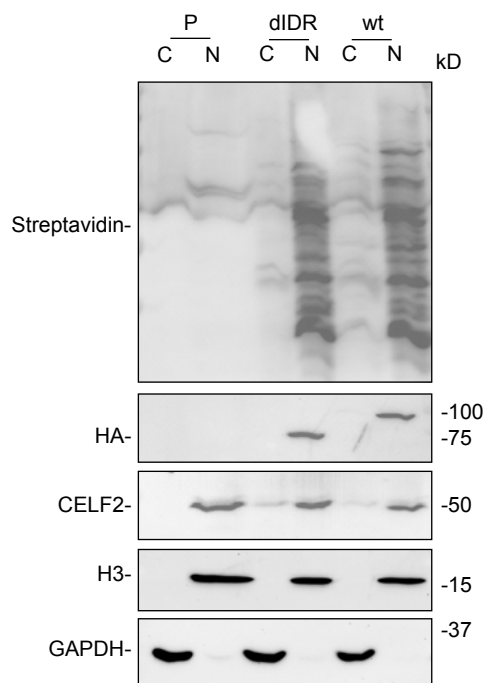
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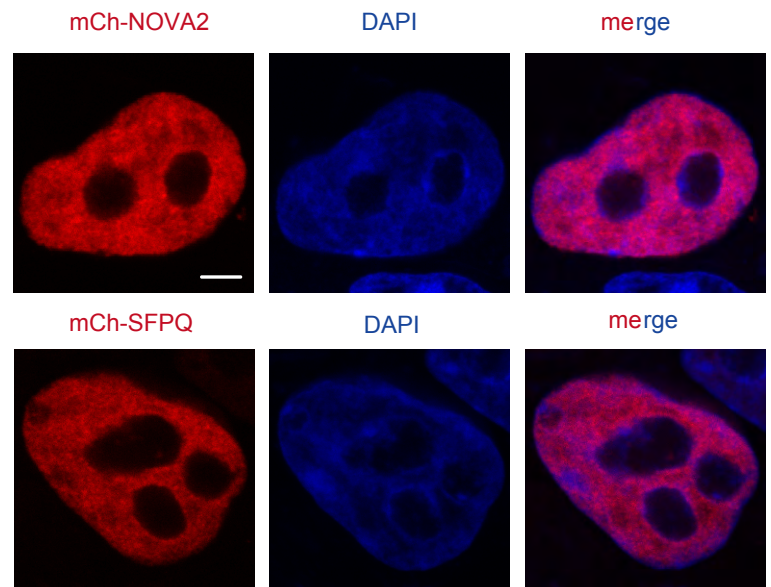
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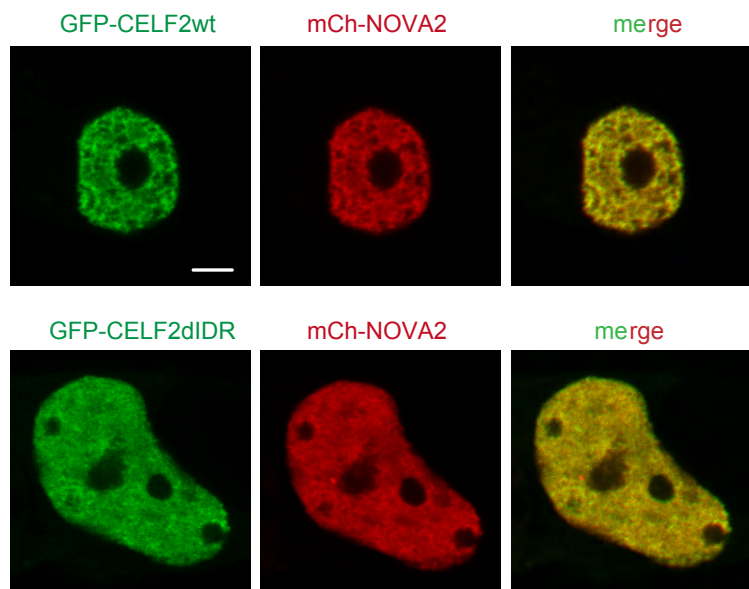
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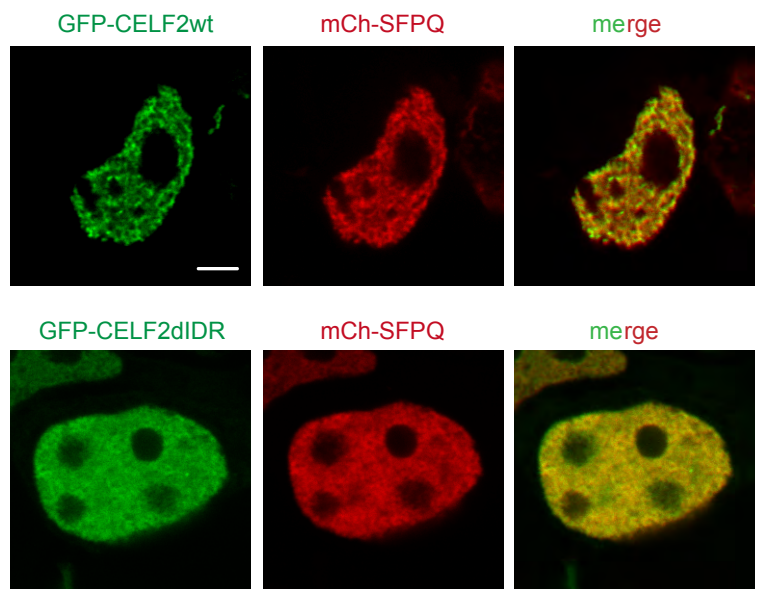
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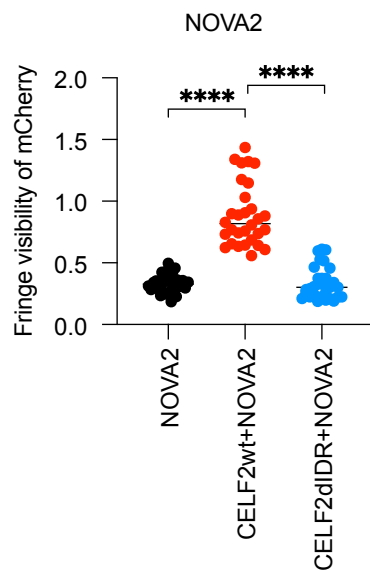
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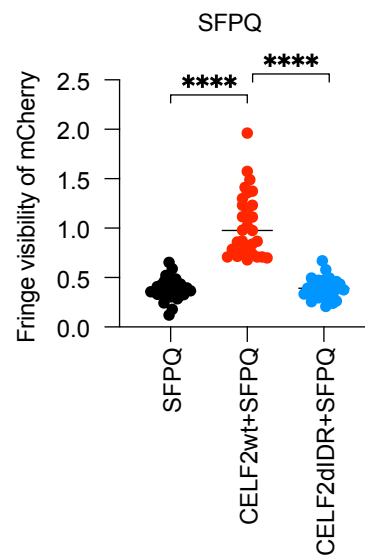
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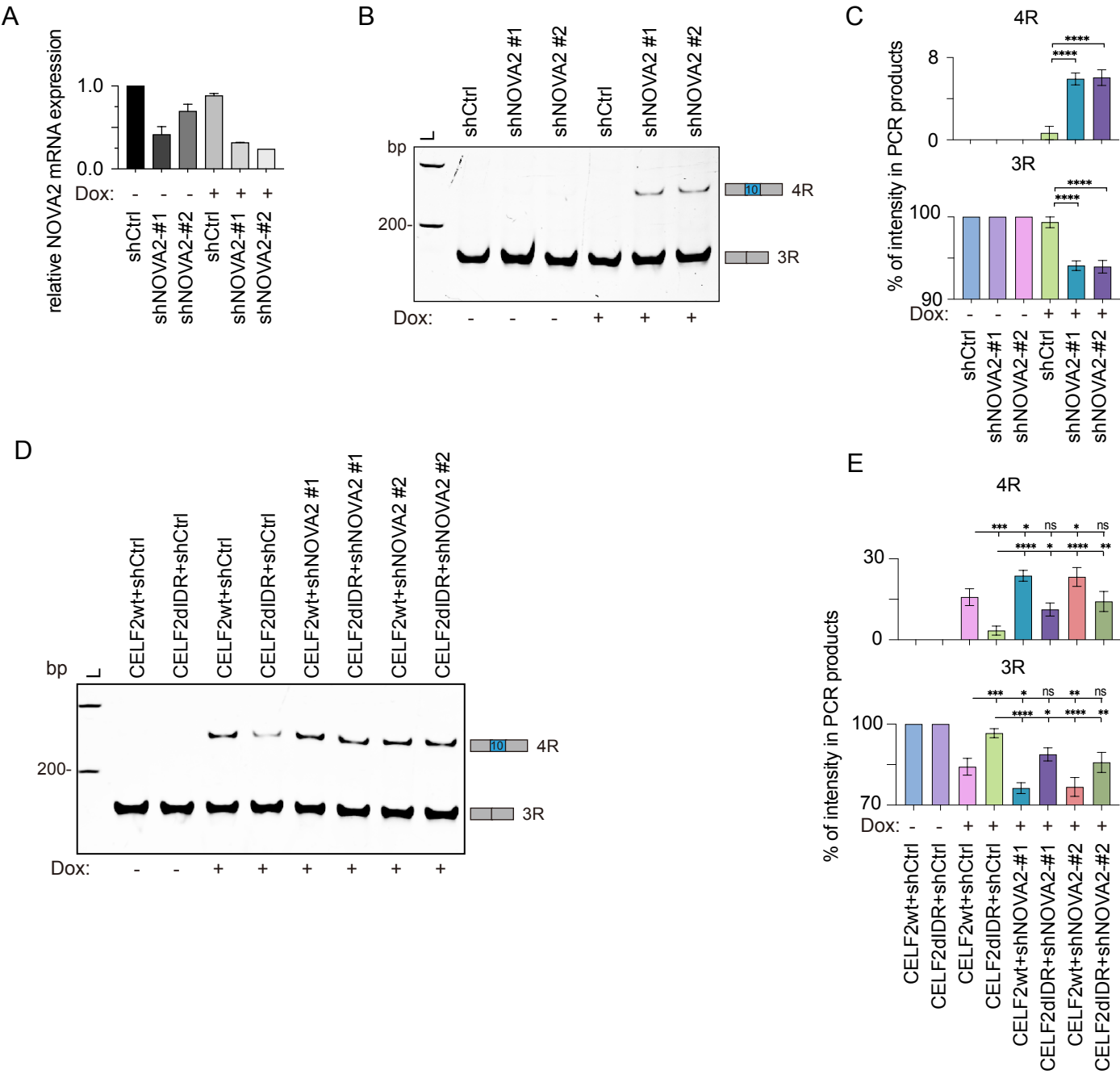


D

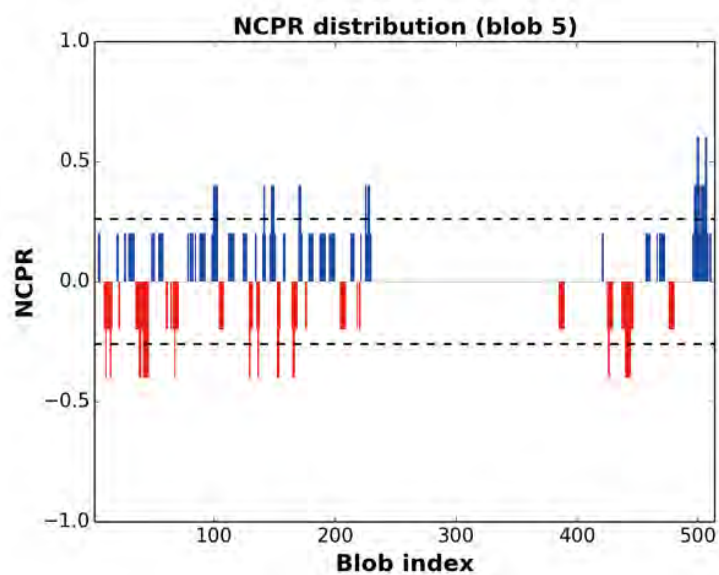


F





A



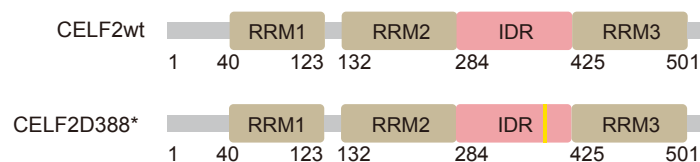
B

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 CELF3 -----
 CELF4 G-----GSTPPGITAPAVPSIPSPIGVNG-----
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 CELF6 A-----ANSPPGSGPCTLPGLPAPIGVNG-----

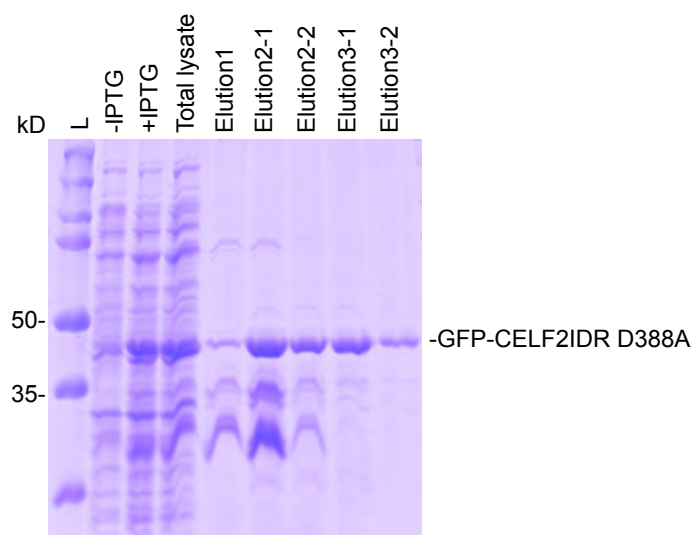
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 CELF2 GLNNINALAVAQ--MLSGMAALNGGLGATGLTNGTA-GTMDALTQAYSGI
 CELF3 -----AQSPAAPVDPLQQAYAGM-----
 CELF4 ---FTGLPPQANGQPAAEAVFANGIHPYP--AQSP-TAADPLQQAYAGV
 CELF5 ---FAGVVPFPGGHPALETVYANGLVYP--AQSP-TVAETLHPAFSGV
 CELF6 ---FGPLTPQTNGQPGSDTLYNNGLSYP--AQSP-GVADPLQQAYAGM

CELFI1 QQYAAAALPT-LYNQN--LLTQQSIGA-----AGSQ
 CELF2 QQYAAAALPT-LYSQS--LLQQQ--SA-----AGSQ
 CELF3 QHYTA-AYPA-AYSLVAPAFPQPPALVAQQPPPPPPQQQQQQQQQQQQQQQ
 CELF4 QQYAGPAYPA-AYGQISQAFPQPPPMI-----PQQQ
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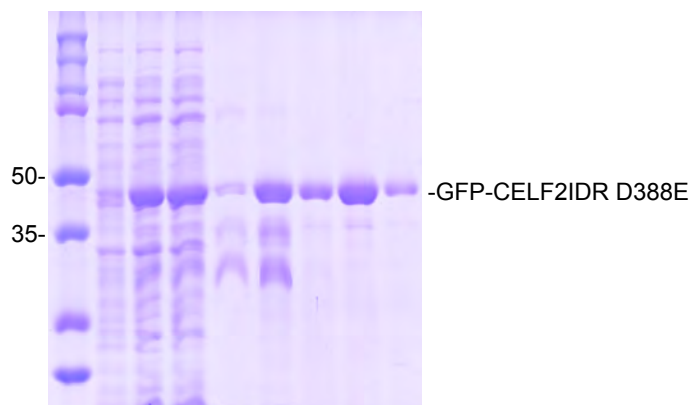
C



D



E



F

