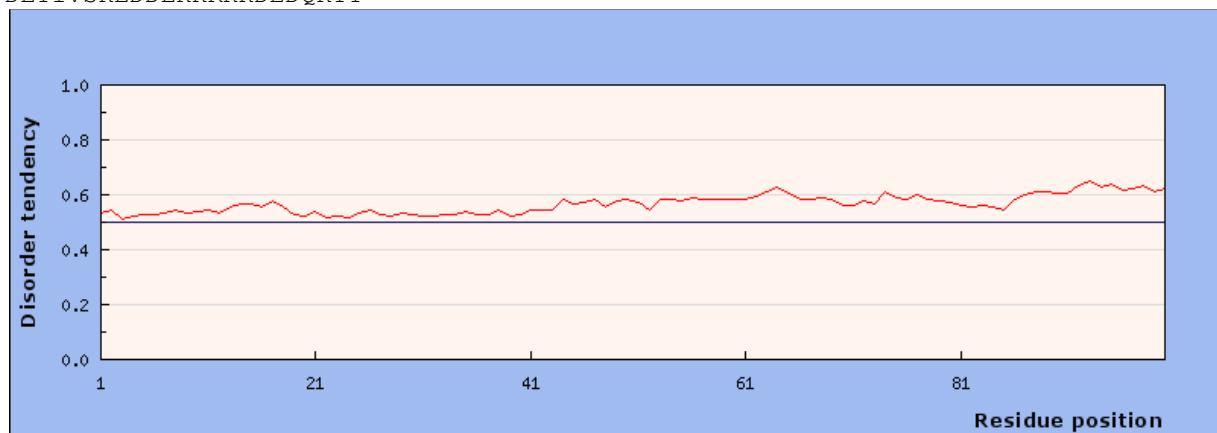


Supplementary Figure S1

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Seq_2

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DT DT QDT KDD RA KTR DD KDD KRG

Seq_3

TKA LET DTY QDD MKG L SMD DTT YQL TTK KET RT KDD TH DDE RSY TLG DD KG KV TR TY DGL KK RQ DF ETE GAK TD
KRD KK STT ITG KG DR T KDK KG TT

Seq_4

RDA EDKRI ARRES RTLY DEQ KSE I DDTR FKT KKE QYG KRE TRED L VRE KR RT QY DR KV QTER I GQ DDL RY RQ TE EDG
YRK TDR K VY DTK D EDD GKV KV

Seq_5

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DRY SER KRD REY KEEN MEKE EIR

Seq_6

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TN DTK SMD TGT DSK DIL TMM TH K

Seq_7

SSK ISE KTT TIS GGAE VY RD KYE SKG TT KAKK KSEE GEY KMD GI KT TO AKV GGT MKT GDG STR IK NY YEE IS KG DYE
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Seq_8

TSM KTK DDK SMD HLKG DTD KTT GH DT KTD NM TLKG DM KG DL DR VRY ET Y TS HD T KIT DG KGT TDD KSV TMN
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Seq_9

NEARR DR RIG D YG DKME ER NR KDE YREAK RG ID YDDR STR LD KQR K RAR D DEDI KTK ED YET VY KES RR ID K DDS I
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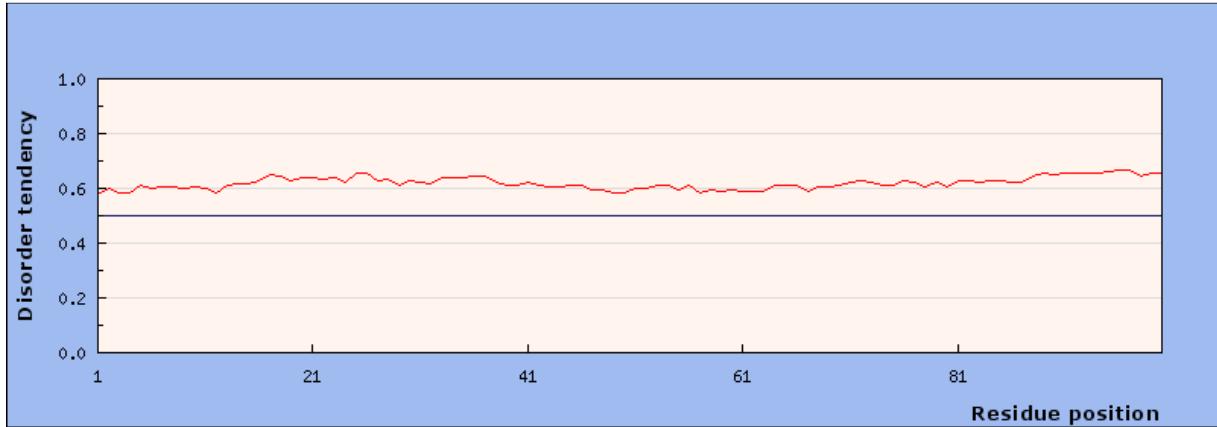
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Dt # 0.6

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Seq_2

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Seq_3

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Seq_4

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Seq_5

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Seq_6

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Seq_7

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Seq_8

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Seq_9

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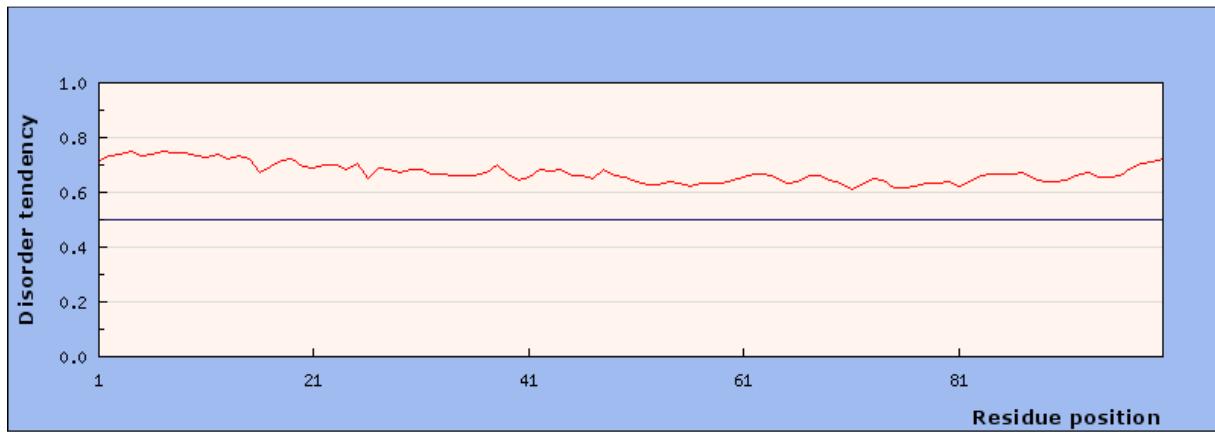
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Dt # 0.65

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Seq_2

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Seq_4

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Seq_5

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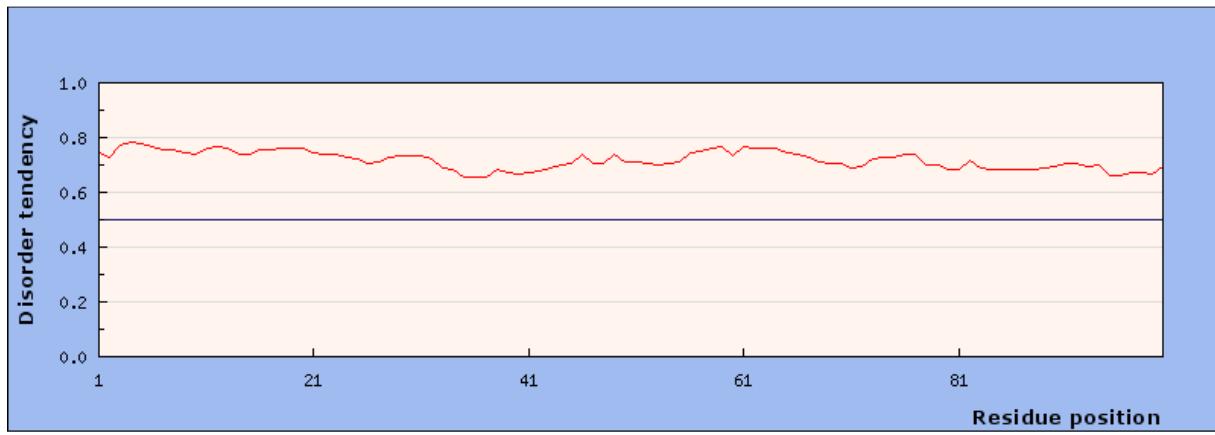
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Dt # 0.7

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Seq_2

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Seq_4

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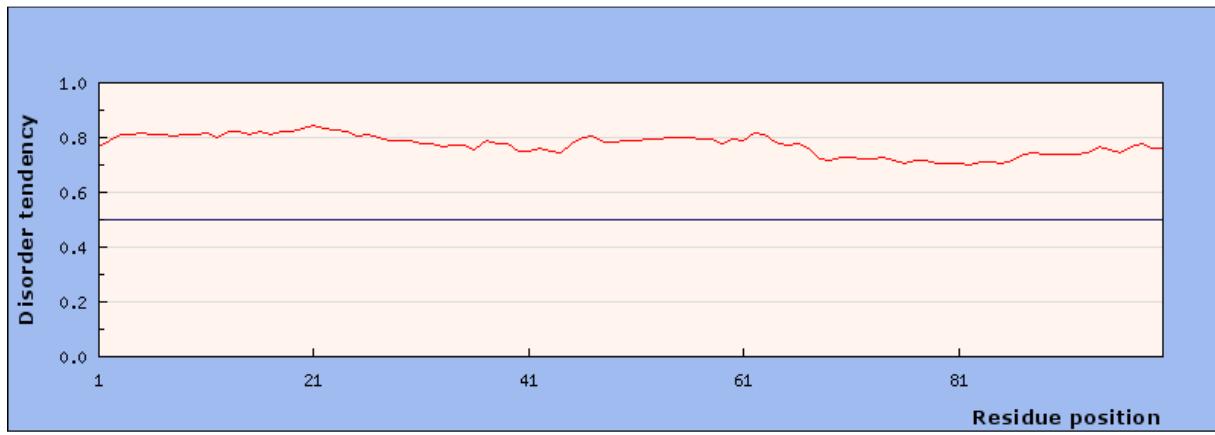
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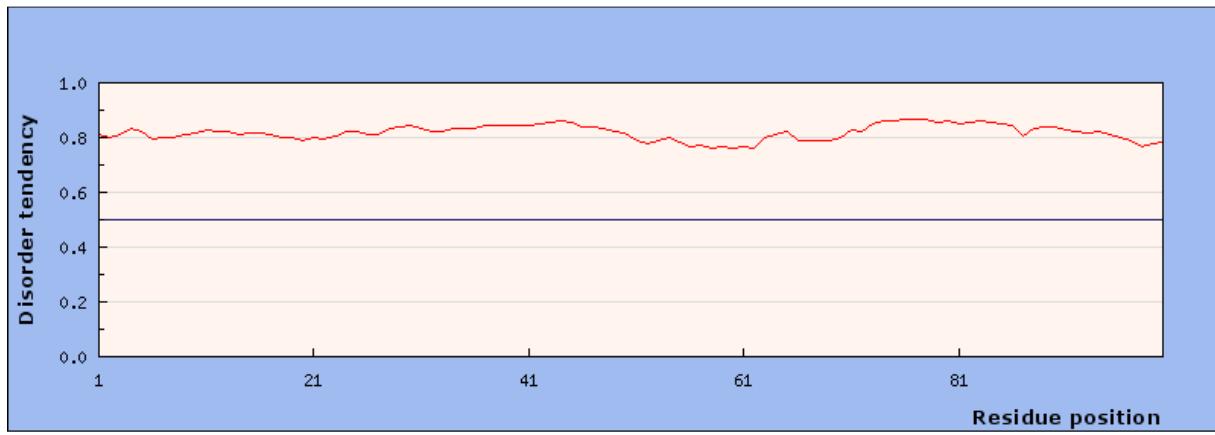
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Seq_3

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Seq_4

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Seq_5

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Seq_7

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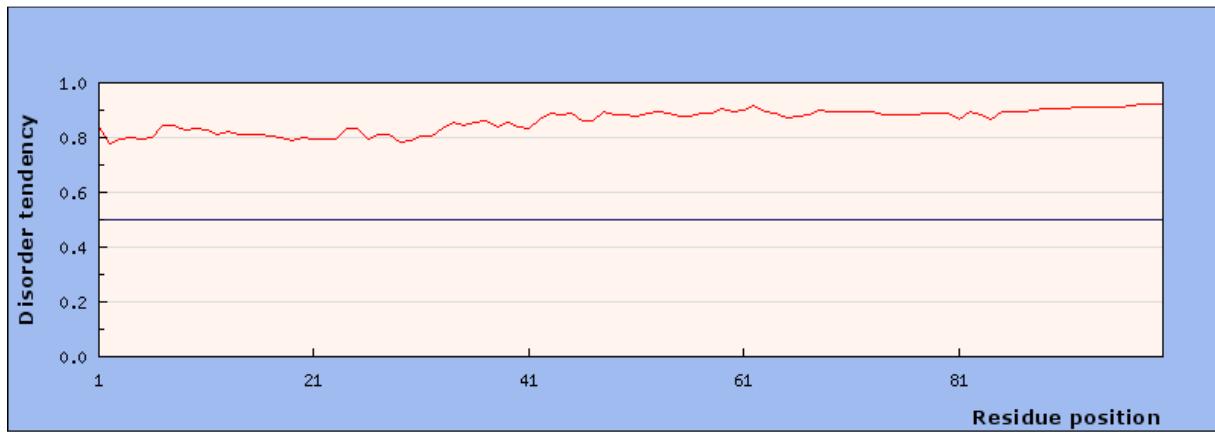
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Seq_2

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Seq_4

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Seq_9

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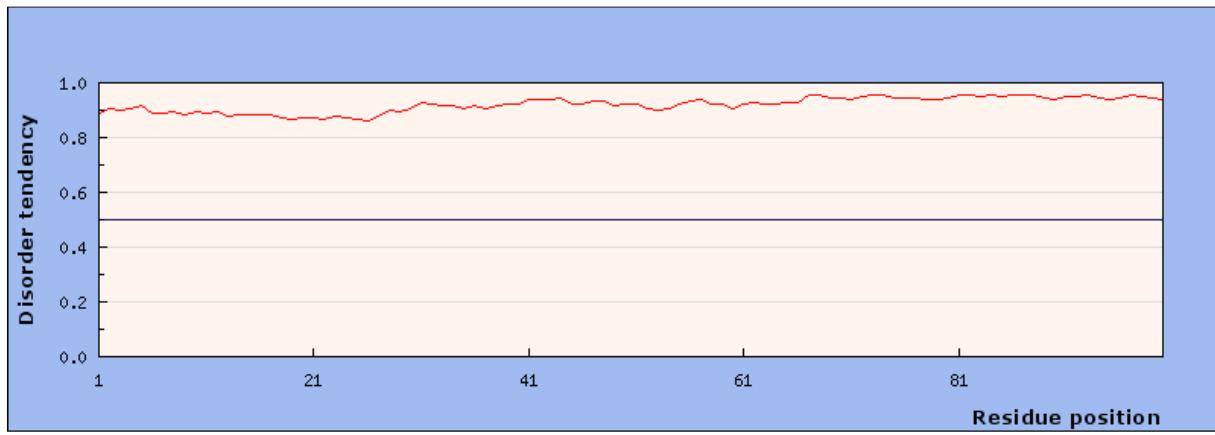
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Seq_2

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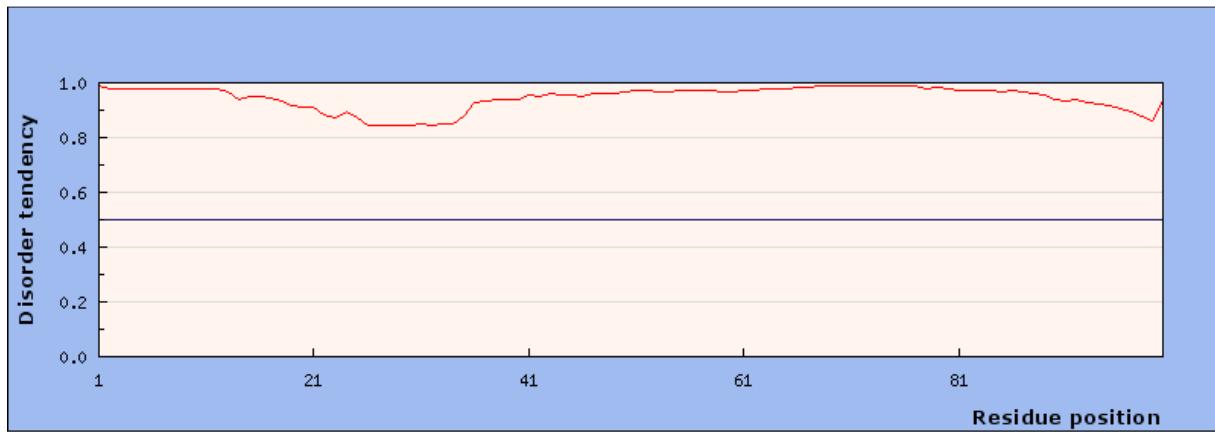
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Dt # 0.95

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Seq_2

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Seq_4

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Seq_5

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Seq_10

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Supplementary Figure S2

NGFP-MoRE

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-----B1MoRE----->

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<-----B2MoRE----->

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P Q D S R R S A D A L L R L Q A M A G I S E - -

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<-----T7 term---->
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NGFP-100-0 . 6MoRE

-----T7 prom----->

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TAA**AG**CG**AC**CG**AA**AA**AG**GT**AC**CC**AG**CA**AA**CG**CG**AG**AC**CC**AC**CA**AA**AC**AG**AC**CC**G**AT**GG**AC**CC**AC**CC**AC**CG**AT**GT**GG**
CG**AC**AA**AC**AG**AC**CC**AC**CG**CG****AT**CC**AA**AG**AT**CC**AA**AG**GT**GG**AT**GC

CCGCAG**GG**AC**AG**TC**GA**AG**GT**C**AG**CT**GAC**CC**CT**G**CT**TA**GG**CT**GCA**AG**CC****AT**GG**CA**GG**AA**AT**CT**CG**GA**AA**TA**ATA**AA**
P Q D S R R S A D A L L R L Q A M A G I S E - -

-----**ACCCAGCTTCTGTACAAAGTGGTGGATCCGGCTGCTAACAAAGCCGAAAGGAAGCTGAGTTGGCTGCT**

<-----T7 term---->
GCCACCG**CT**GA**GC**AA**TA**ACT**AG**