

Table of tested synthetic Notch receptors

Constructs	AA sequence	Containing Domains
FNQ1	886-1772 (Fly)	10EGF, NRR, TM, PEST
FNQ2	1142-1772 (Fly)	5EGF,NRR, TM, PEST
FNQ3	1259-1772 (Fly)	3EGF, NRR, TM, PEST
FNQ4	1452-1772 (Fly)	NRR, TM, PEST
FNQ5	886-1772 (Fly)	10EGF, NRR, TM
FNQ6	1142-1772 (Fly)	5EGF, NRR, TM
FNQ7	1259-1772 (Fly)	3EGF, NRR, TM
FNQ8	1452-1806 (Fly)	NRR, TM
FNQ9	1452-1772 (Fly)	NRR, TM
MNQ	1427-1753 (Mouse)	NRR, TM
WNQ1	20-794 (Worm)	5EGF, NRR, TM, PEST
WNQ2	480-794 (Worm)	NRR, TM, PEST
WNQ3	480-794 (Worm)	NRR, TM

DNA Sequences used in the study.

Sequence of GBN:

GCCGATGTGCAGCTGGTGGAGTCTGGGGGAGCCTTGGTGCAGCCGGGGGGGTCTCTGAGACTCTCCTGTGCAGC
CTCTGGATTCCCCGTCAATCGCTATAGTATGGGGTGGTACCGCCAGGCTCCAGGGAAGGAGCGCGAGTGGGTGCG
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CGCCAGGAATACGGTGTATCTGCAAATGAACAGCCTGAAACCTGAGGACACGGCCGTGTATTACTGTAATGTCAA
TGTGGGCTTTGAGTACTGGGGCCAGGGGACCCAGGTCACCGTCTCCTCA

Sequence of QF2.0 with 3XMyC tag:

CCACCCAAGCGCAAAACGCTTAACGCTGCGGCTGAGGCTAACGCTCATGCCGACGGACACGCCGACGGAAACGCC
GACGGACACGTGGCCAATACGGCCGCGTCCTCGAATAATGCGAGGTTGCTGATCTCACTAACATCGATACTCCG
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GGAGGCGGTGGCTCCGAACAGAACTTATCAGCGAGGAGGATCTCGGCGAGCAGAAGCTGATTTCCGAAGAAG
ACCTTGAGAGCAAAAGCTCATATCCGAAGAGGACCTGTCTG

Sequence of SP-GFP-mcd8 (GFP sequence in green color):

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Sequence of Serrate cytosol domain:

AAACAGCGTCTGGCTTATCGCACCAGTTCGGGAATGAACTTAACTCCCTCGCTGGATGCACTGCGTCACGAGGAG
GAGAAGTCGAATAATCTGCAGAACGAGGAGAATCTGCGAAGGTATACAAATCCGCTGAAGGGCAGCACCAGTTC
CCTAAGAGCGGCCACCGGCATGGAATAAGCCTCAATCCCGCTCCGGAATTAGCCGCCTCGGCGGCGAGTAGTTC
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GCAGGCGACAAGCGGAGCTCACAGATTCTGCTGCACAAAACCCAAAACCTCGGACATGCGGAAGAACACTGTGG
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Sequence of DI cytosol domain:

TCGGTGGCGGATAACAACAATGCCAACTCAGACTTTTGTGTGGCTCCGCTACAAAGAGCCAAGTCGCAAAAGCAA
CTCAACACCGATCCACGCTCATGCACCGCGGTTCCCGGCAGGCAGCTCAGCCAAGGGAGCGTCTGGCGGAGG

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GGCGGCGGCGGGAGTGGCCGGAGCCTGTTTCATCCCACTAATGGCTGCAGCTTCGGTAGCGGGCAGCGGAGCG
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Sequence of LDL endocytosis signal:

GGTAACTTCGCTAACCCCGTGACGAGTCCATGTAC

GPI anchoring sequence:

ACCAGCAATAGCCGCACCAGCACCGATAAGGGAGAGCCCAACACCGGCGGCGACCGTTACATCTATGGCCATACC
TGCCTGATTACCCTCACCGTCCTGCACGTTATGCTGTCCTTGATCGGCTACTTGACCAACTCCGCTACTTCCAGCTCC
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Fly Notch 10XEGF+NRR+TM domain:

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Fly Notch 5XEGF+NRR+TM domain:

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Fly Notch 3XEGF+NRR+TM domain:

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Fly Notch NRR+TM domain:

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Fly Notch PEST domain:

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Mouse Notch NRR+TM domain:

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Gfwd21

Worm Notch 5EGF+NRR+TM domain:

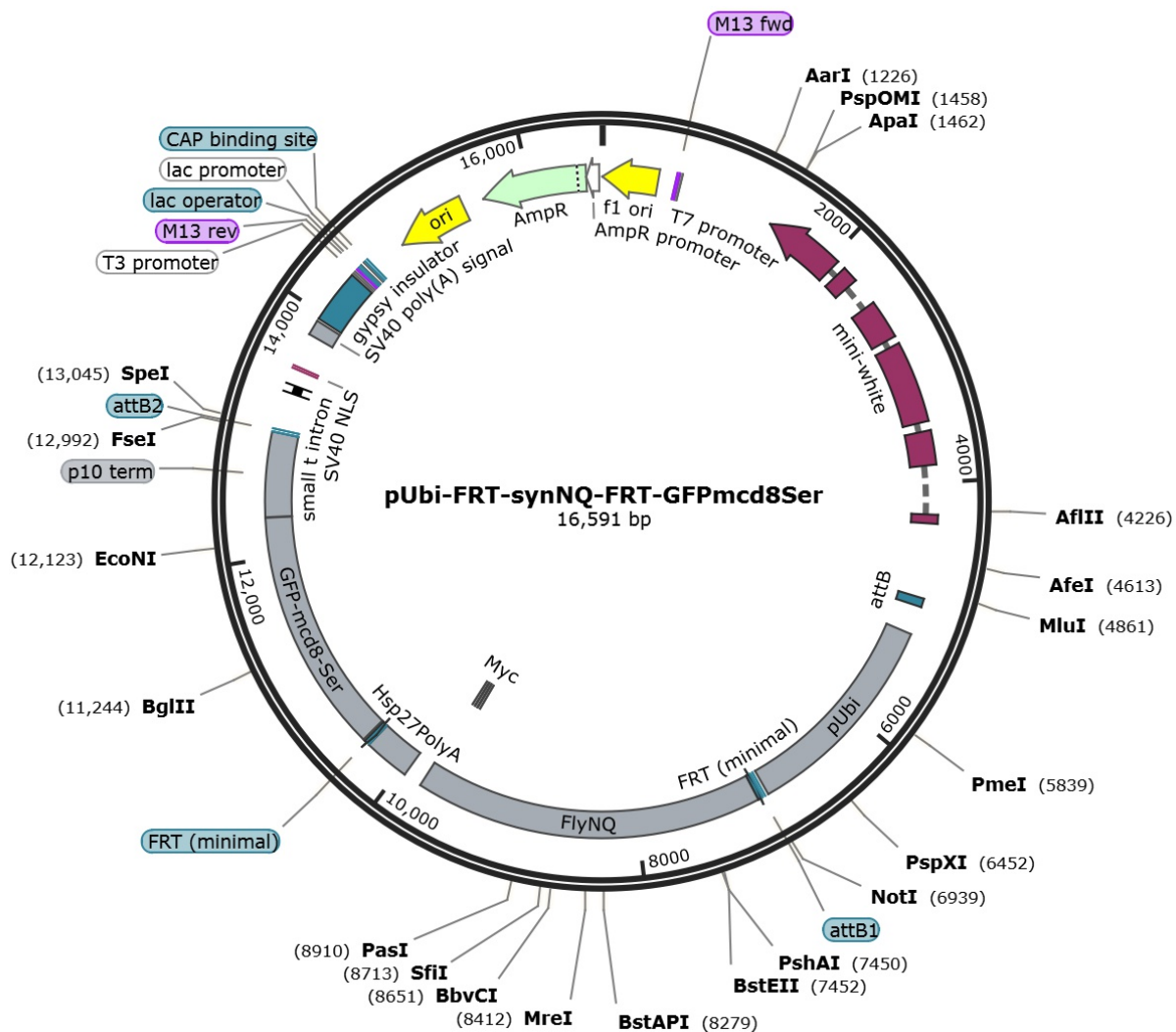
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Worm Notch NRR+TM domain:

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Full Sequence of *pUbi-FRT>synNQ.Stop.FRT>GFPmcd8Ser*



Sequence of the vector in Genbank format:

LOCUS Exported 16591 bp ds-DNA circular SYN 15-MAR-2017

DEFINITION synthetic circular DNA

ACCESSION .

VERSION .

KEYWORDS pUbiFRTflyQFRTmcd8

SOURCE synthetic DNA construct

ORGANISM synthetic DNA construct

REFERENCE 1 (bases 1 to 16591)

AUTHORS .

TITLE Direct Submission

JOURNAL Exported Wednesday, Mar 15, 2017 from SnapGene Viewer 3.3.2

<http://www.snapgene.com>

FEATURES Location/Qualifiers

source 1..16591

/organism="synthetic DNA construct"

/mol_type="other DNA"

rep_origin complement(6..461)

/direction=LEFT

/note="f1 ori"

/note="f1 bacteriophage origin of replication; arrow
indicates direction of (+) strand synthesis"

primer_bind 603..619

/note="M13 fwd"

/note="common sequencing primer, one of multiple similar
variants"

promoter 626..644

/note="T7 promoter"

/note="promoter for bacteriophage T7 RNA polymerase"

CDS complement(join(1444..2058,2129..2260,2481..2796,2858..3512,
3587..3860,4255..4326))

/codon_start=1

/gene="white"

/product="Drosophila white gene eye color pigment"

/note="mini-white"

/note="This is a modified version of the white gene lacking
part of the first intron."

/translation="MGQEDQELLIRGGSKHPSAEHLNNGDSGAASQSCINQGFGQAKNY"

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protein_bind 4923..4992

/bound_moiety="phage phi-C31 integrase"

/note="attB"

/note="attB site for the phi-C31 integrase (Groth et al.,
2000)"

misc_feature 5219..6929

/note="pUbi"

protein_bind 6956..6980

/gene="mutant version of attB"

/bound_moiety="BP Clonase(TM)"

/note="attB1"

/note="recombination site for the Gateway(R) BP reaction"

protein_bind complement(6983..7016)

/bound_moiety="FLP recombinase from the Saccharomyces
cerevisiae 2u plasmid"

/note="FRT (minimal)"

/note="supports FLP-mediated excision but not integration
(Turan and Bode, 2011)"

misc_feature 7018..9798

/note="FlyNQ"

CDS 9697..9726

/codon_start=1

/product="Myc (human c-Myc oncogene) epitope tag"

/note="Myc"

/translation="EQKLISEEDL"

CDS 9730..9759

/codon_start=1

/product="Myc (human c-Myc oncogene) epitope tag"

/note="Myc"

/translation="EQKLISEEDL"

CDS 9763..9792

/codon_start=1

/product="Myc (human c-Myc oncogene) epitope tag"

/note="Myc"

/translation="EQKLISEEDL"

misc_feature 9946..10310

/note="Hsp27PolyA"

protein_bind complement(10322..10355)

/bound_moiety="FLP recombinase from the Saccharomyces

cerevisiae 2u plasmid"

/note="FRT (minimal)"

/note="supports FLP-mediated excision but not integration

(Turan and Bode, 2011)"

misc_feature 10371..12314

/note="GFP-mcd8-Ser"

misc_feature 12319..12994

/note="p10 term"

protein_bind complement(12996..13020)
/gene="mutant version of attB"
/bound_moiety="BP Clonase(TM)"
/note="attB2"
/note="recombination site for the Gateway(R) BP reaction"

intron 13327..13392
/note="small t intron"
/note="simian virus 40 (SV40) small t antigen intron"

CDS 13522..13542
/codon_start=1
/product="nuclear localization signal of SV40 large T antigen"
/note="SV40 NLS"
/translation="PKKKRKV"

polyA_signal 13814..13935
/note="SV40 poly(A) signal"
/note="SV40 polyadenylation signal"

protein_bind 13957..14387
/bound_moiety="Su(Hw), CP190, and Mod(mdg4)67.2"
/note="gypsy insulator"
/note="chromatin insulator from Drosophila"

promoter complement(14406..14424)
/note="T3 promoter"
/note="promoter for bacteriophage T3 RNA polymerase"

primer_bind complement(14445..14461)
/note="M13 rev"
/note="common sequencing primer, one of multiple similar variants"

protein_bind 14469..14485

/bound_moiety="lac repressor encoded by lacI"

/note="lac operator"

/note="The lac repressor binds to the lac operator to inhibit transcription in E. coli. This inhibition can be relieved by adding lactose or isopropyl-beta-D-thiogalactopyranoside (IPTG)."

promoter complement(14493..14523)

/note="lac promoter"

/note="promoter for the E. coli lac operon"

protein_bind 14538..14559

/bound_moiety="E. coli catabolite activator protein"

/note="CAP binding site"

/note="CAP binding activates transcription in the presence of cAMP."

rep_origin complement(14847..15435)

/direction=LEFT

/note="ori"

/note="high-copy-number ColE1/pMB1/pBR322/pUC origin of replication"

CDS complement(15606..16466)

/codon_start=1

/gene="bla"

/product="beta-lactamase"

/note="AmpR"

/note="confers resistance to ampicillin, carbenicillin, and related antibiotics"

/translation="MSIQHFRVALIPFFAAFCCLPVFAHPETLVKVKDAEDQLGARVGYI
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LIKHW"

promoter complement(16467..16571)

/gene="bla"

/note="AmpR promoter"

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