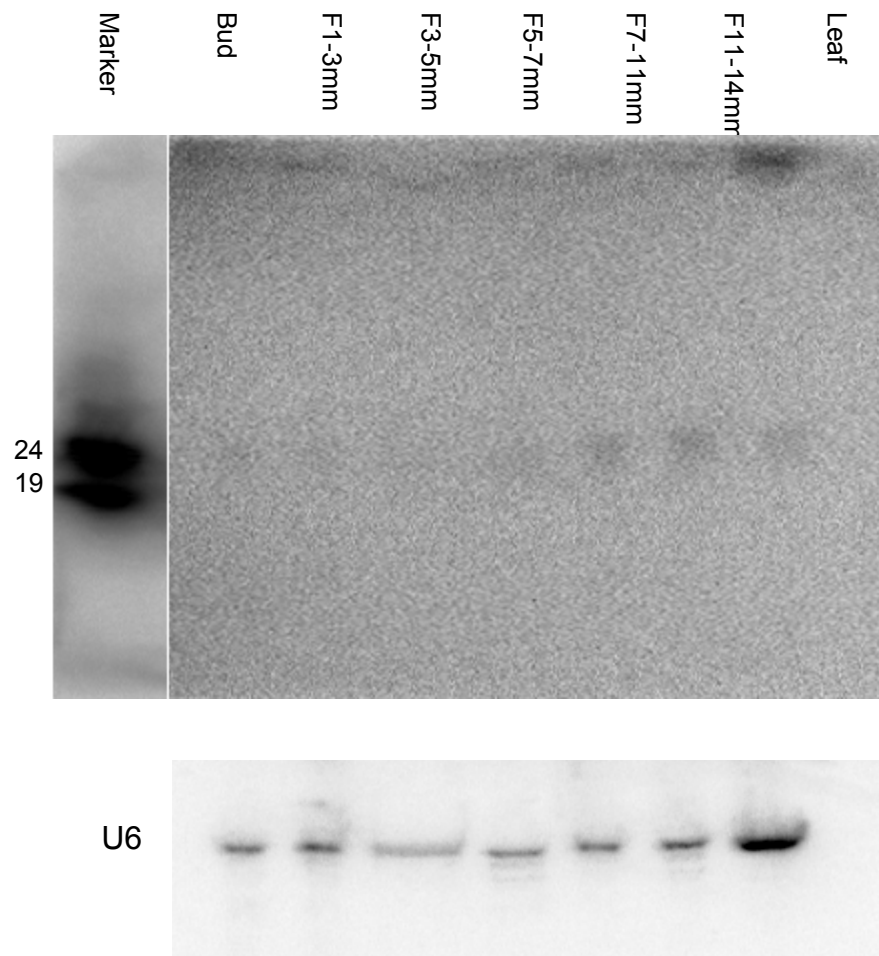


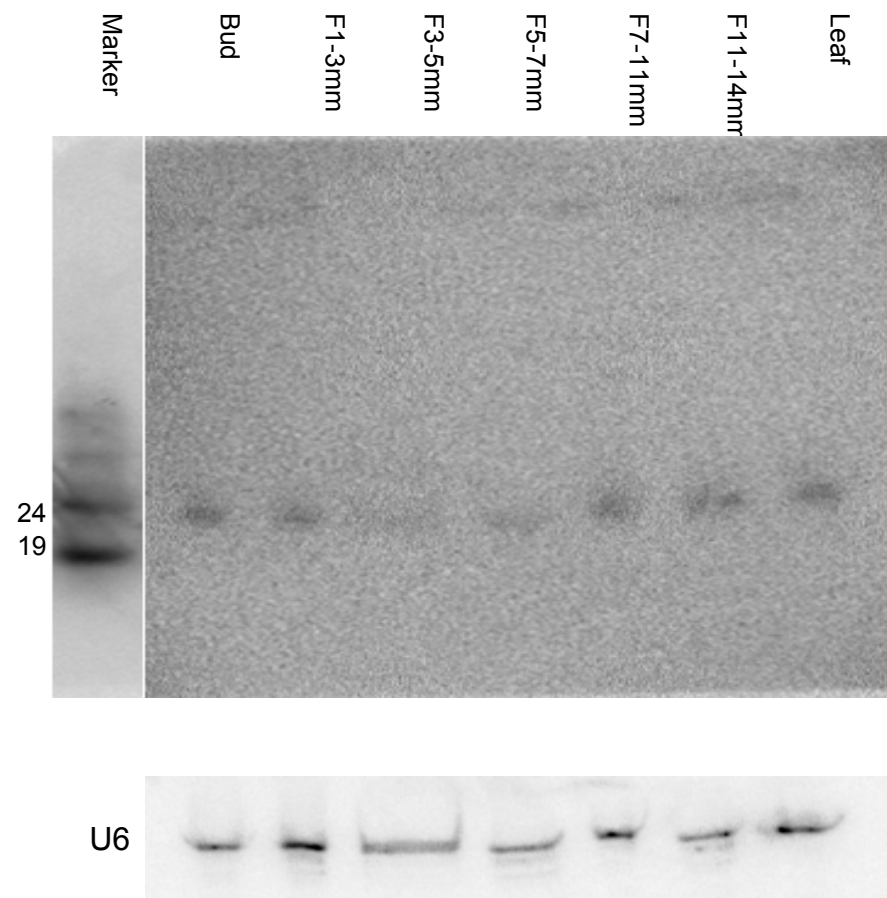
Oligo name	Sequence 5'-3'
Tom-192	TTCGGACTCTTCAAAAATATC
Tom-23	AGAAGTCAAACCTCGCTTCTTG
Tom-204	ACCCACGGGATAGTGTCACGT
Tom-117	TAAGTGTCTATTTATGCACAC
Tom-105	CATGGCACACTACATGACAATTT
Tom-208	TATACTATACCAGGGTCTATTA
Tom-30	GTTTGATATCGTAACAACAATT
Tom-127	GACACAATCTTATCCATCGATA
Tom-185	AGATGTGCACAAGTAGACACTT
Tom-5	AGTTTAGGGTCCAATATATATA
Tom-7	TACTGTGAGTAACATATATCAAT
Tom-9	GTCACCTTTTATTAAAGTCAC
Tom-10	TGAAAAAGTCACAATCTTTCAG
Tom-50	TCTGAGTGAACCAGGCCAATAT
Tom-66	AATATATATATTGCTTGCACCAA
Tom-76	AATAACTATATTTTTAGGATTAT
Tom-80	AACGTATTCCACAATAACGTTT
Tom-87	GAACTTTCACATATAGCCACT
Tom-92	AACAACATTAGGTTGATATACGTT
Tom-95	CAAGTTTAACTGTCTACTTGTGC
Tom107	TAGTAACGTATATCAACCTAATG
Tom-110	ACACACATGTCTTACATGACATAA
Tom-122	CCGCGCCTTTCTCACTTATAC
Tom-123	TAAATAAACACATGAGTCTTATAT
Tom-126	CCTCGAGCCAGACAACATTCC
Tom-129	GTCTGAATCAGAATCATTGAGATT
Tom-141	CATGACACCCTACATGACAA
Tom-144	AACACATGAGTCTTACATGACAC
Tom-149	TACATGTATTAGGTCATGTAG
NGM1	ATCCAGCACACGGTAGCTTCG
NGM2	TCACCAAGATCTGCACCGA
NGM3	ATCCAGCATCATCTCTCCTGCG
NGM4	CAGCACACGGTAGCTTCGCGCCA
NGM5	GTCATAATCCAGCACACGGTA
NGM6	CAGTCATAATCCAGCACACGG
Tom-54	TGACGTGACCGAATTATAATTGG
Tom-157	AACGTTACCGTGATATACGTTT
Tom-161	CCCTACATGAGGTGTCTAAATGA
Tom-169	AAACTAGTATAAACATATTTT
Tom-170	CACTTTTGTTGTATAAAGCG
Tom-173	TGGGGCCTAACTTCCTAACCATT
Tom-174	TAGCAGCTTAATTAATTAATTA
Tom-176	TCATGTAGGACACGTGTGTCTATT
Tom-177	AGACACACGTGTCTTACATGACA
Tom-178	TCGACCTACATCTACAAAGGAG
Tom-179	TTAACATGGTTTCGGTCAATCGAC
Tom-180	TCTCCTTTATGGACGTAAGTCG
Tom-182	TTGGTCTCATTTTCATGTTTAT
Tom-183	ATCTACACCATCCGTTAAATC
Tom-186	GCACCAATGTCCCTAAAGTAT
Tom-193	CCAATTCTCTAGCAAATACAG
Tom-32	GACGAATGTGTCAATTTGTTT
Tom-40	GAAACGTGGCACCCGATCCAT
Tom-44	GGCACGACGGACCGTCACACATCT

Oligo name	Sequence 5'-3'
Tom-52	CTTCAAATTTTAACTTTGACA
Tom-189	TTCCATGGATTGTCTTTGGATA
Tom-191	TAGAGTGACGCAGATCTTAC
Tom-64	TAAATAAACACATGAGTCATACAT
Tom-72	CATGAGTCGTACATAACACA
Tom-82	TATCAATCTGACACCATCAAC
Tom-143	ATAATTAACCAGAGCCAGCTAA
Tom-153	GATATATATGTGCCACGTGGAC
Tom-101	TTTCGTCATACTTTTGGGACATT
Tom-108	GGGTATGCACAAGTAGACACTT
Tom-119	TGCCACGTAGACTAATAAACT
Tom-131	GTTGAAGGGCATAAATGTAAT
Tom-136	ATTAATCAAAACGGCAAAAT
Tom-139	TCGTAGATACGACGGACCGTCA
Tom-154	TTTTGAAAGGGTTGCAACCT
Tom-156	ACGGTAACGTTTTAGGATATATTT
Tom-160	TTGAGGTGTCTAAGTGAAAT
Tom-175	GATGTGCATAAGTAGACACTTA
Top-1	GGATCTCTTCTAGGATAGGCT
Top-2	AGTGGCACCTGTGTCTCCAGA
Top-3	TTCATAAATAAGTGGCACCTGT
Top-4	CAGTCATAATCCAGCACACGG
Top-5	ACATATATAGTCACACCGATTA
Top-6	TGGTACATGAGGCATACTCGCATGT
Top-7	TGACTTAGAGGCGTTTCAGTCAT
Top-8	TCCAGACATGTTGTCTAACCG
Top-9	TAGTCACACCGATTACAGTTCTG
Top-10	TGTGCTTGGGGTAGCCAG
Top-11	GTCGGTTAGGACAACGTGTCTG
Top-12	TGAGACAGTTTGGTTATTTACA
Top-13	GGATCTCTTCTAAGATAG
Top-15	GCCAGATCATGAATTAGTAACT
TomTar1	TAATATATTTCAAAAACAATTAT
TomTar2	GTTTCATTTTAGACACCTCATGTAG
TomTar3	GTTGGTCGTGAAATATCAATG
TomTar4	CAGACACCTGTTGCACCAAT
TomTar5	GAACCGAACCGAATTATTTTCA
TomTar6	GACATGTGACACAATCTTATCTGT
TomTar7	TCTGAATCTTAATCATTGAGATT
TomTar8	CAGGTCAACAGTTCGAGCCCC
TomTar9	CGCACCGGATGTTTACATCAA
TomTar10	GGCCCCCTACTGCGGGTCGGCA
TomTar11	TTCATTGACCACTAGACCAC
TomTar12	GACACCAATGTCCCAAAAGTATG
TomTar13	TCCTACCAAGCACACCCCTAAAT
TomTar14	CCATTAACCCAACTCAATAAA
TomTar15	TTCAATCATATTGCACAAAAA
TomTar16	AAGACACATGTATCTACTTGTTC
TomTar17	TGACCCAAAACCTCATACTTAAC
TomTar18	TTTTCACTTGGACACCTCAAG
TomTar19	TTAAAACAGACAAACAAAT
TomTar20	TTACCCCATCAGGCCCTGCAT

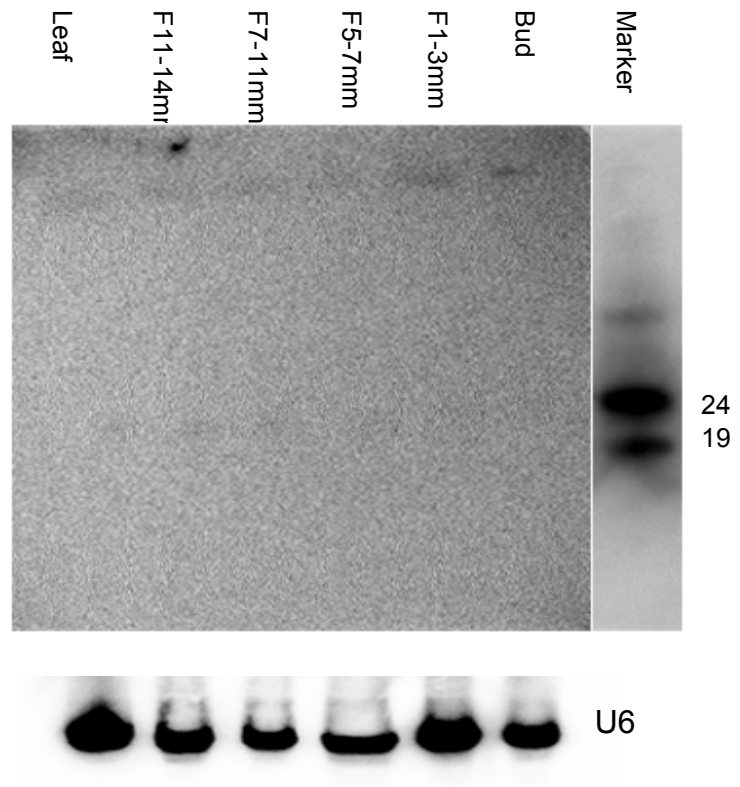
Tom23



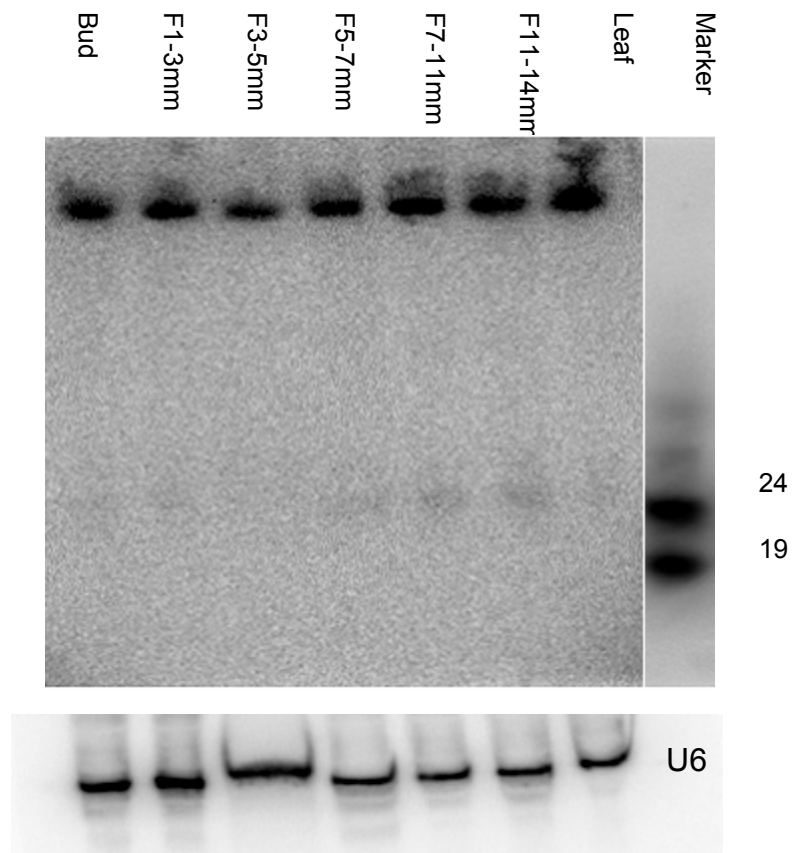
Tom204



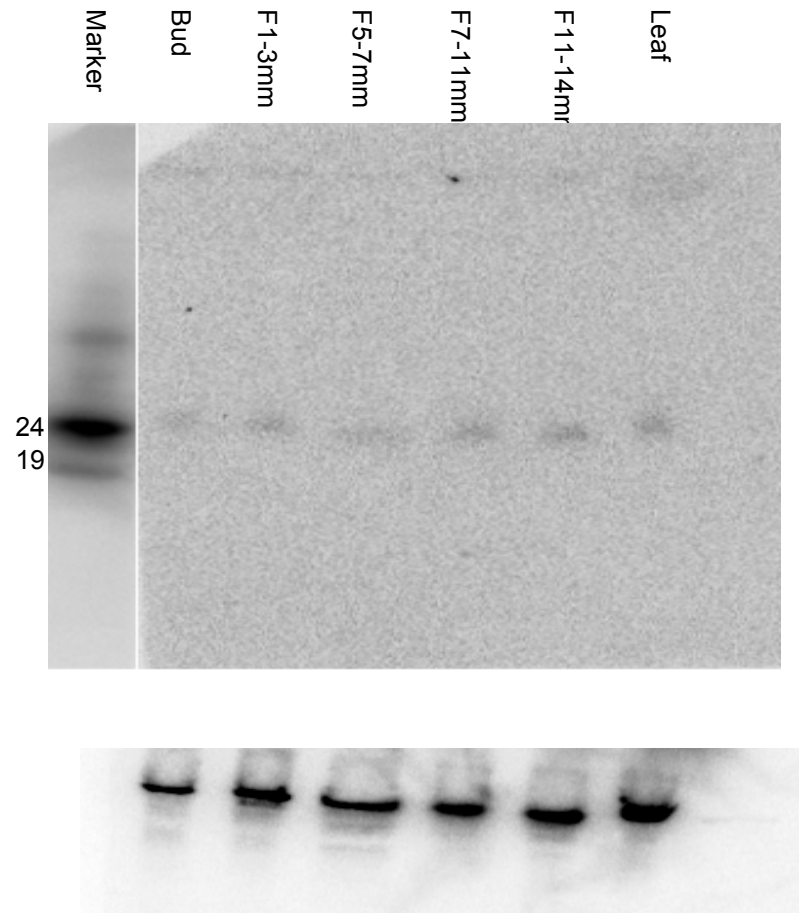
Tom117



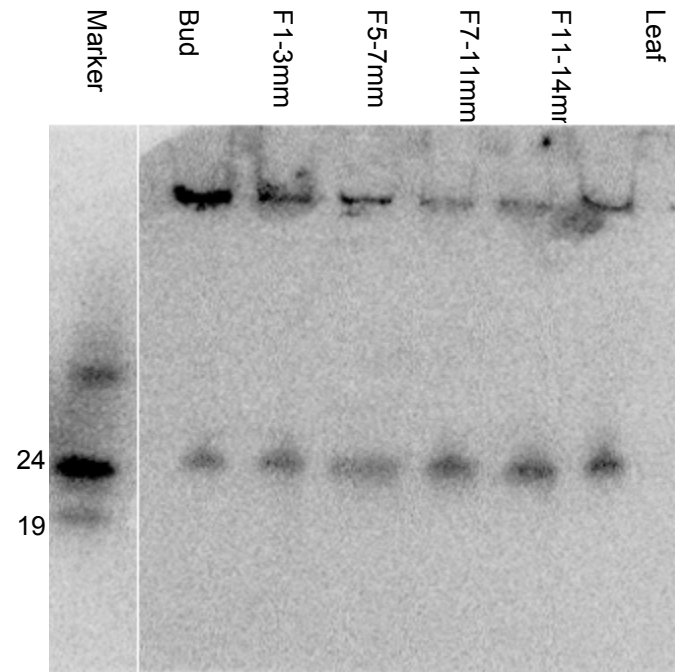
Tom208



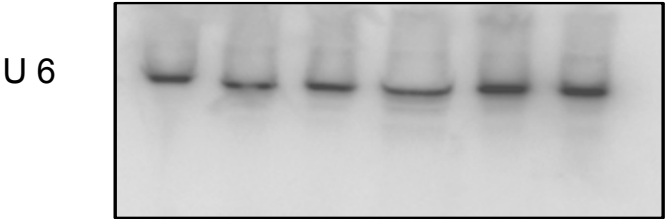
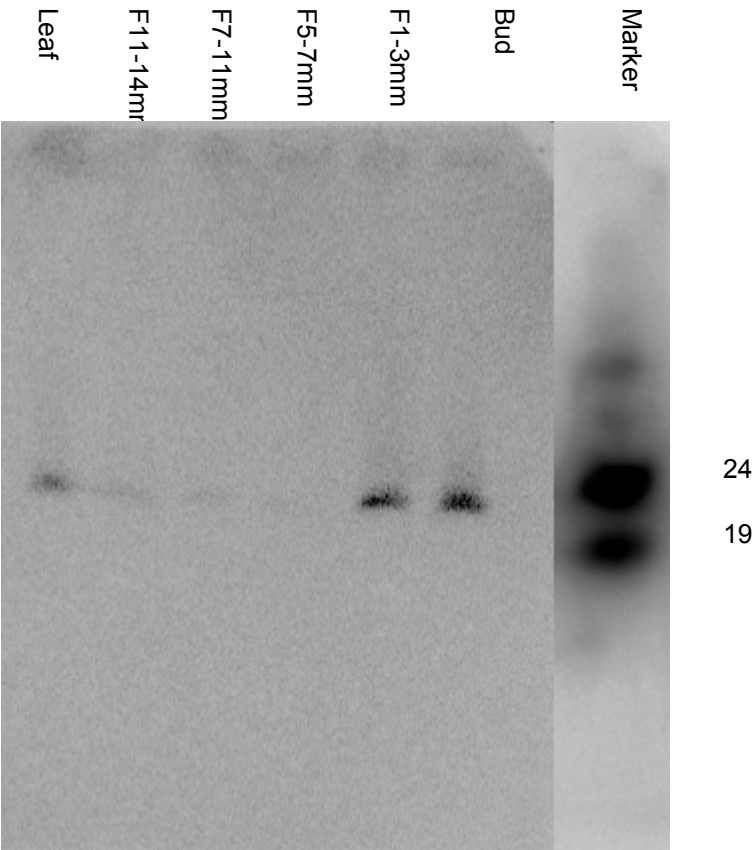
Tom127



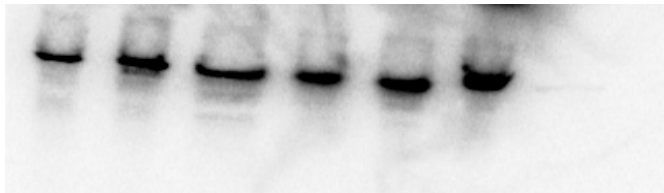
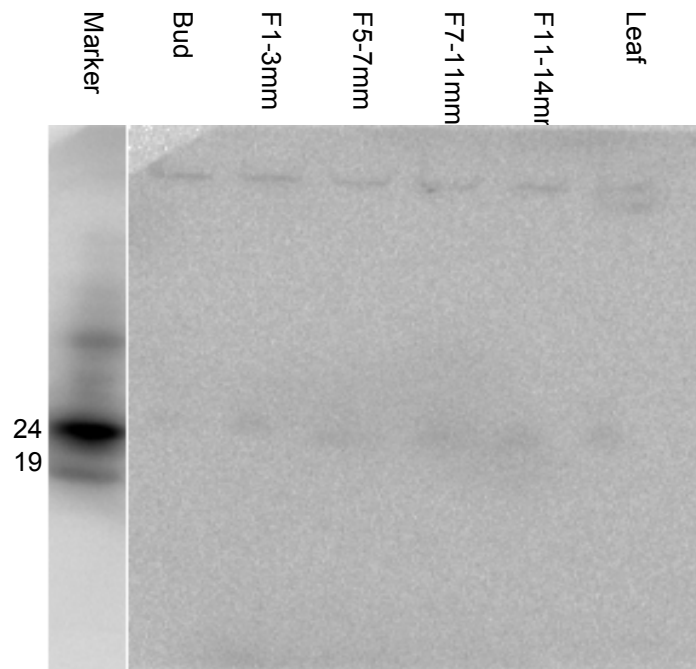
Tom185



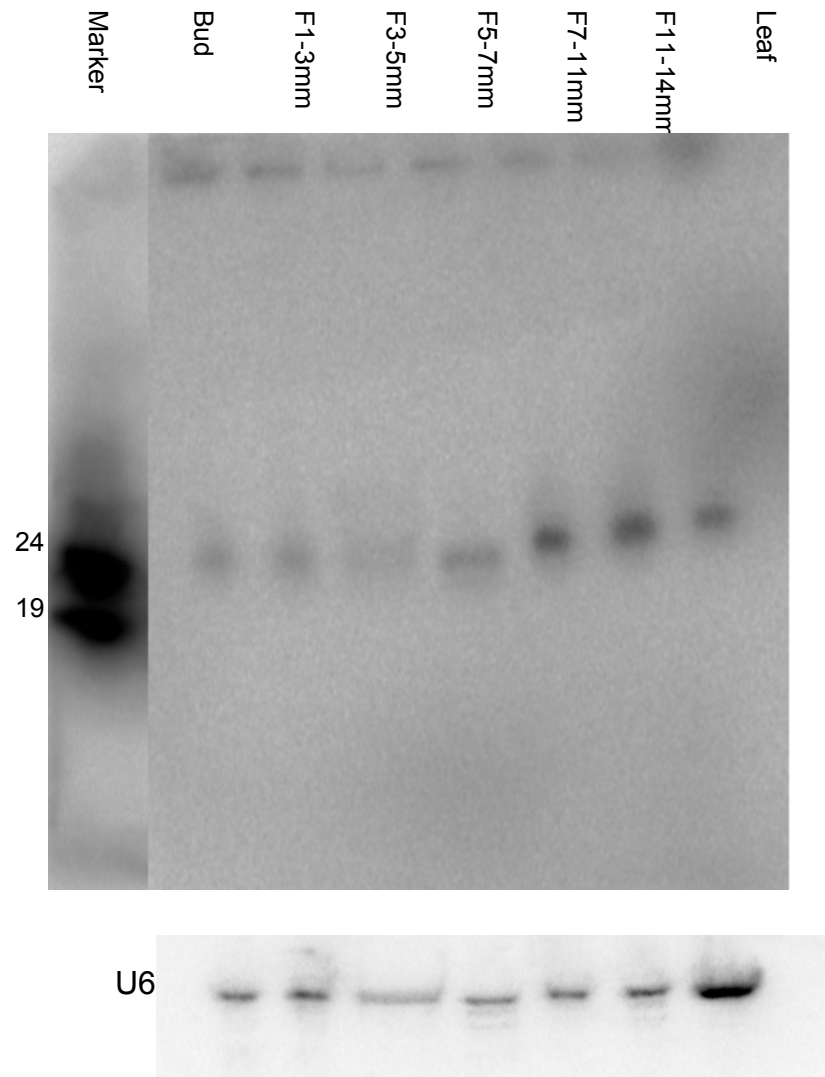
Tom50



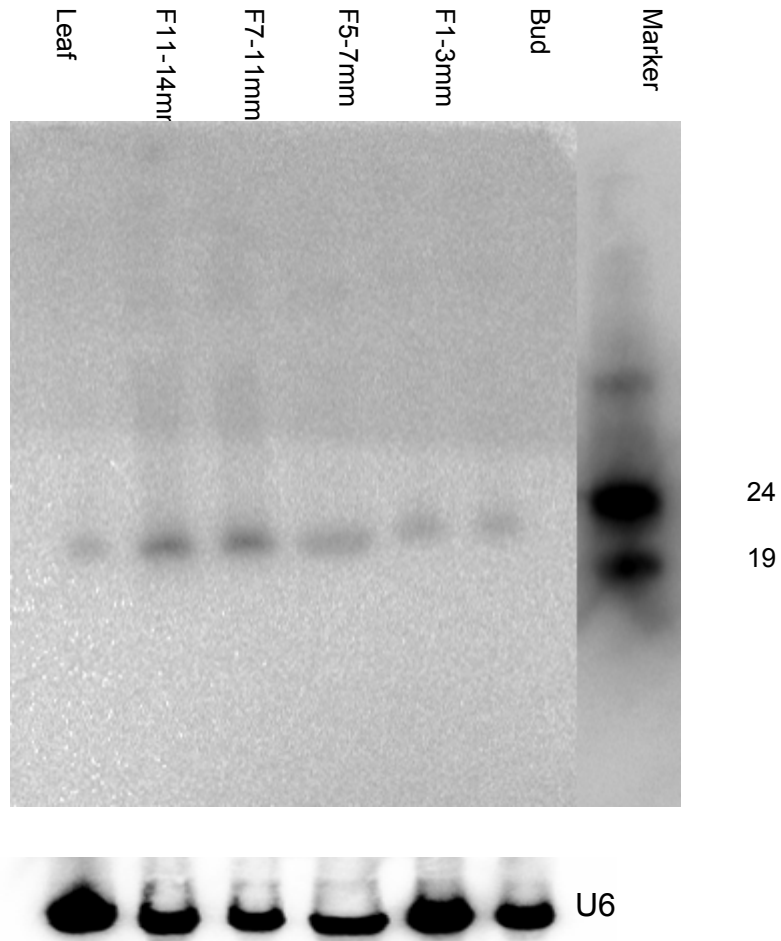
Tom87



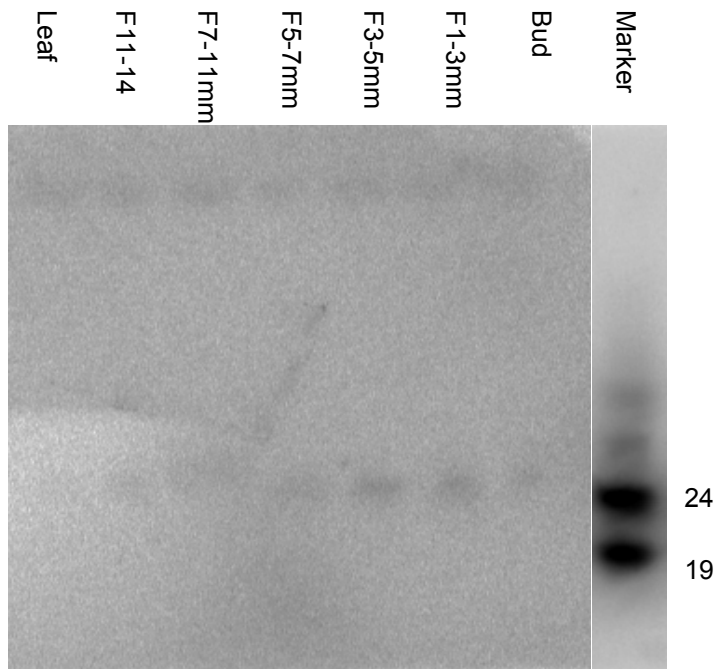
Tom122



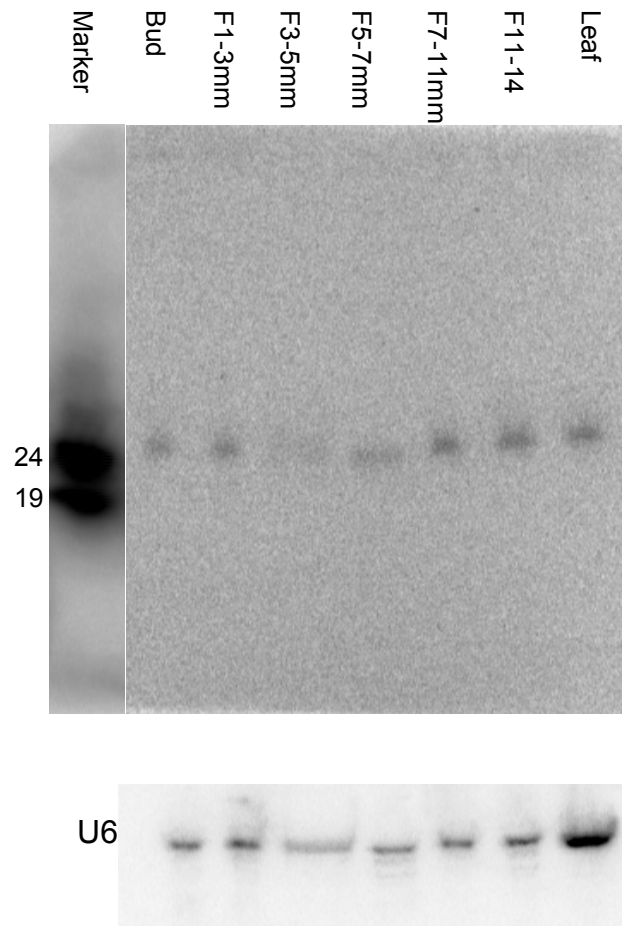
Tom126



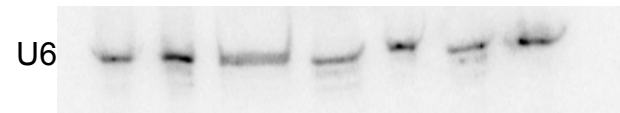
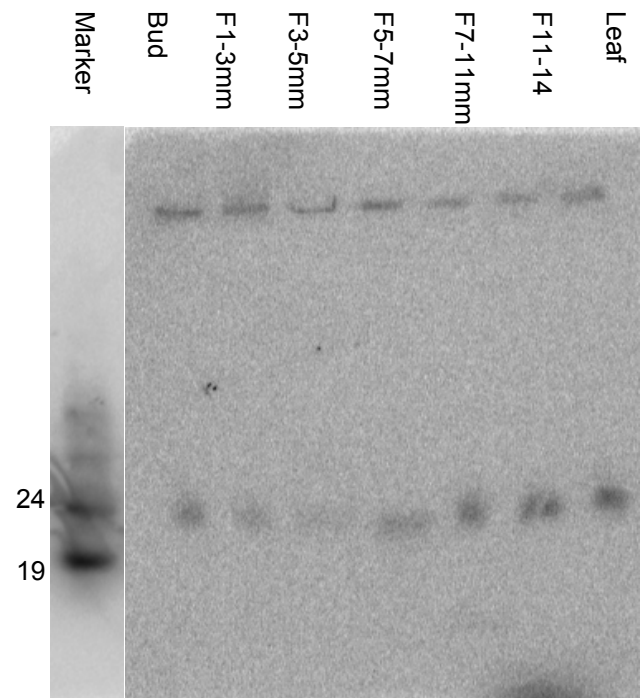
Tom129



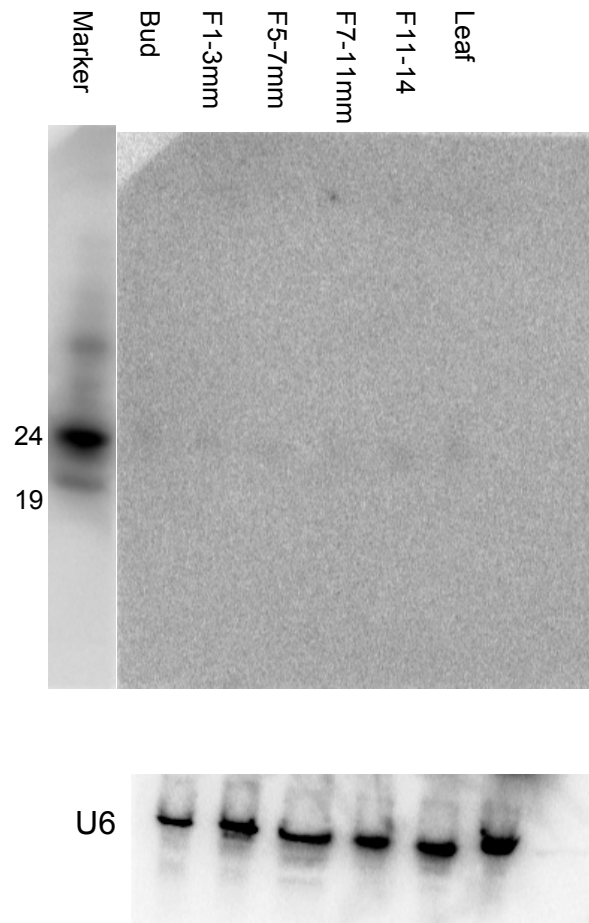
Tom141



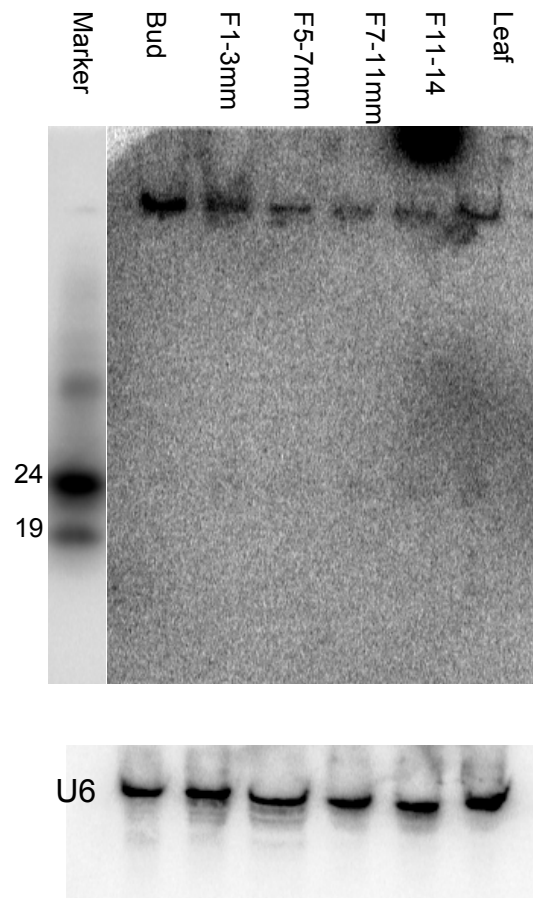
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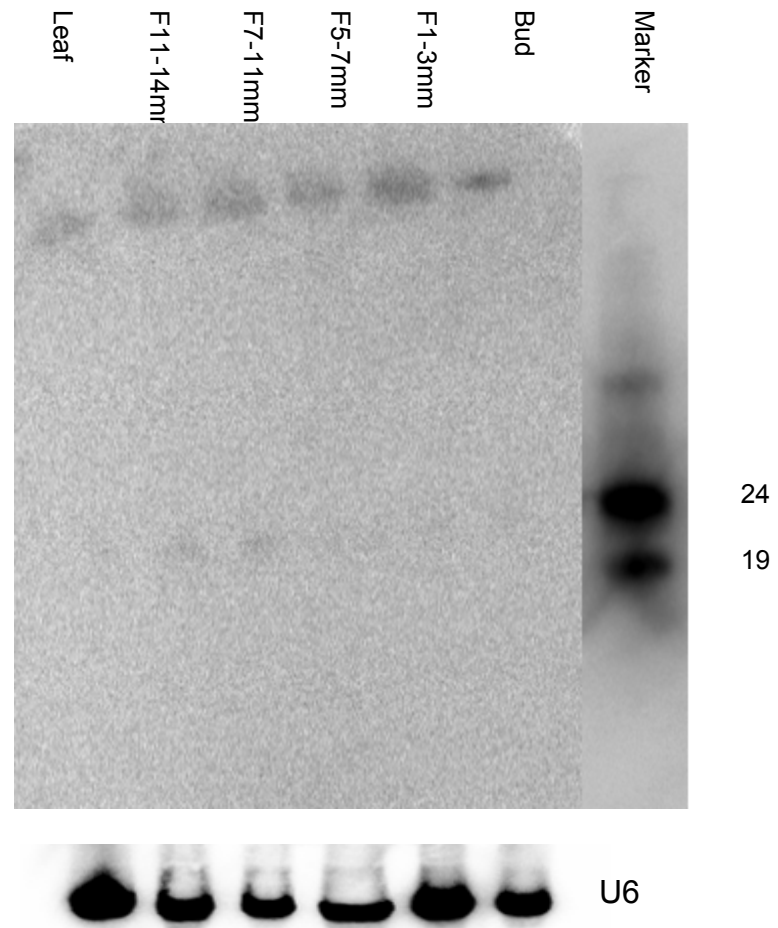
Tom149



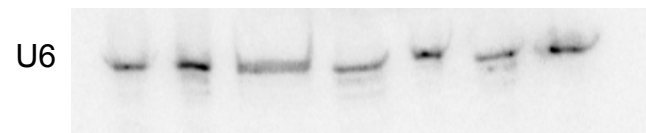
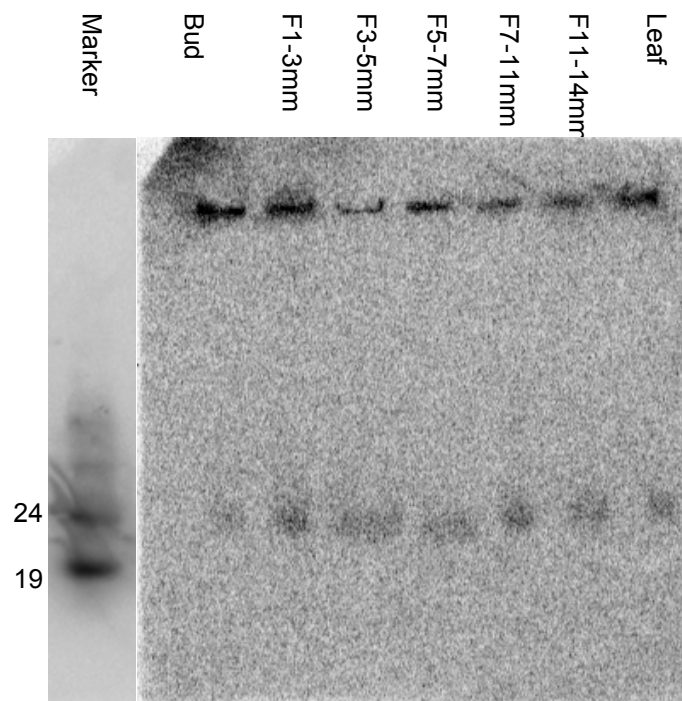
Tom157



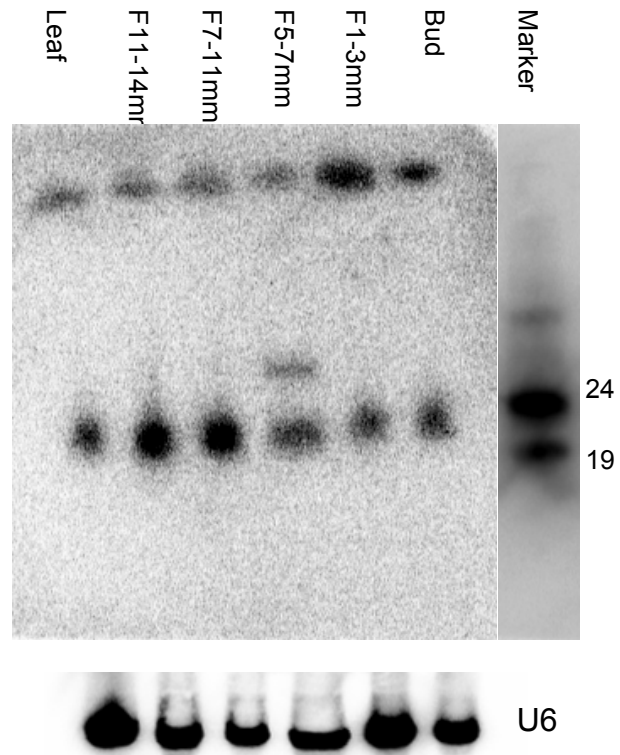
Tom161



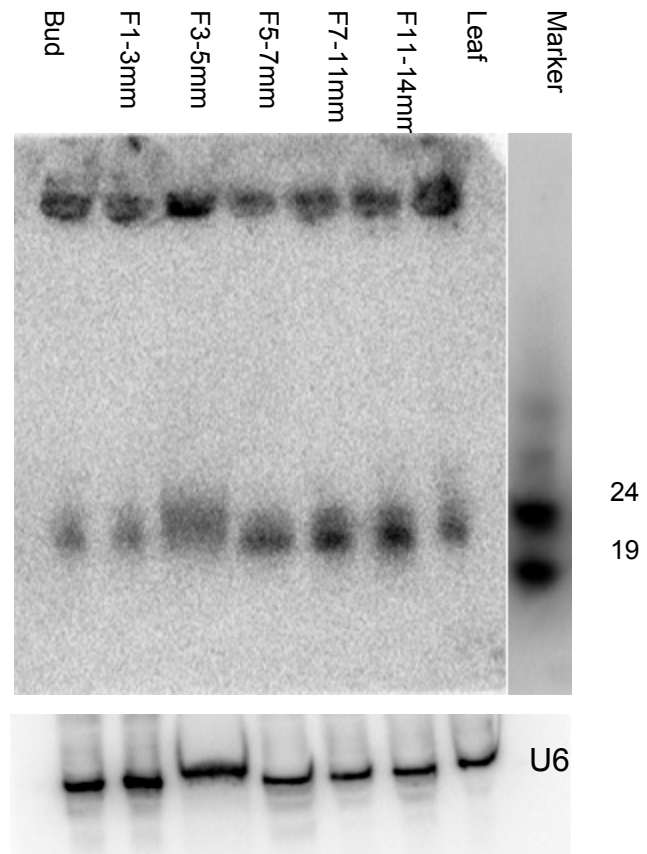
Tom173



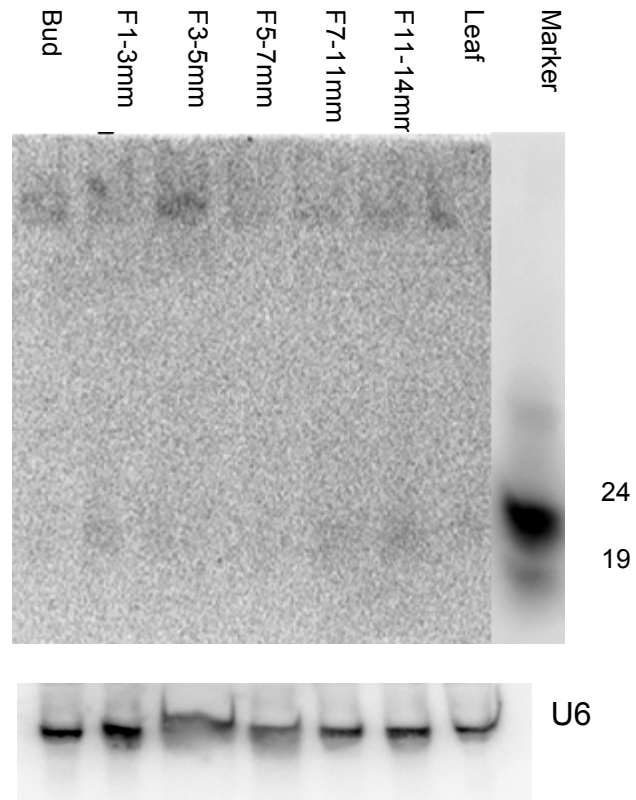
Tom177



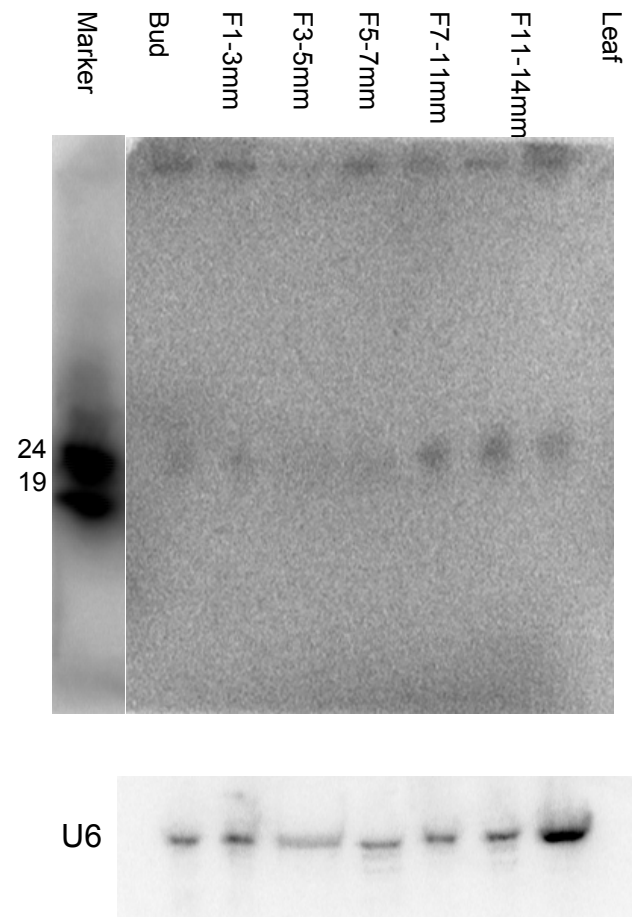
Tom179



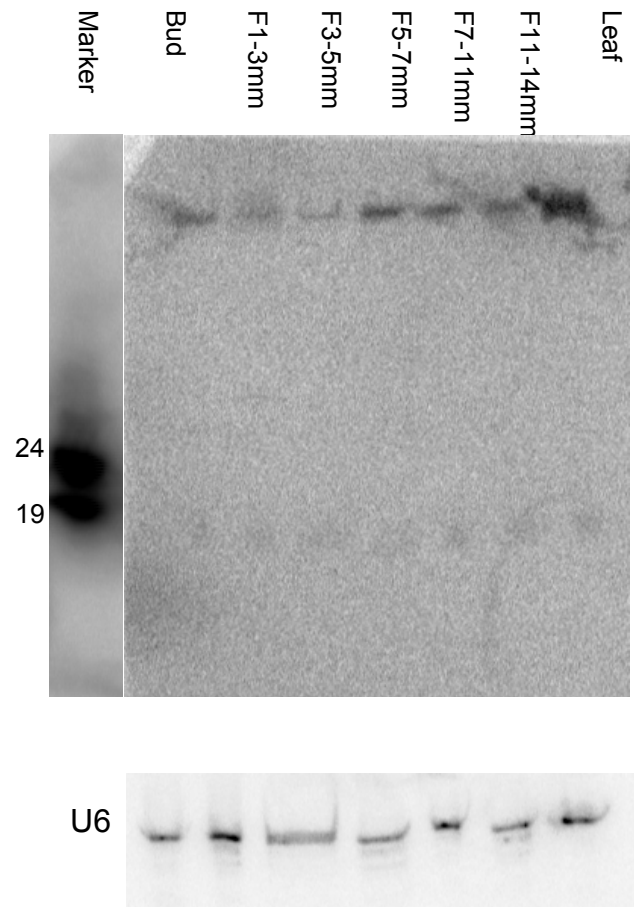
Tom183



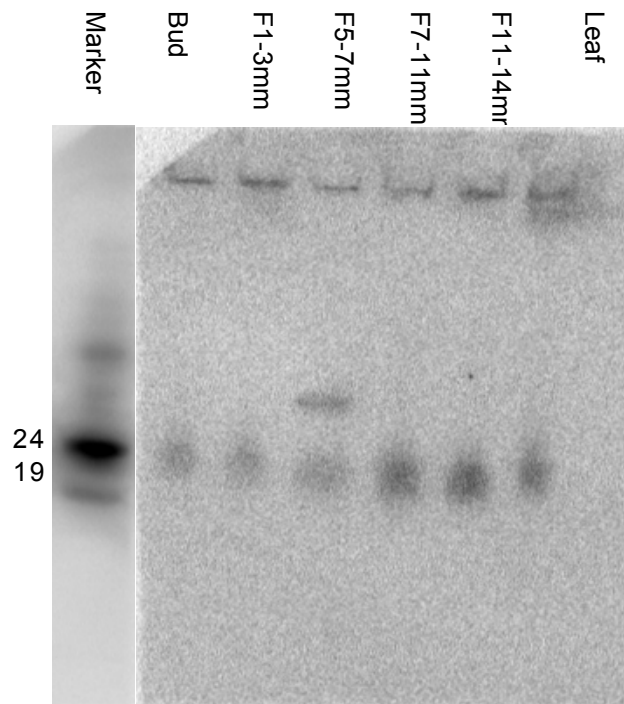
Tom186



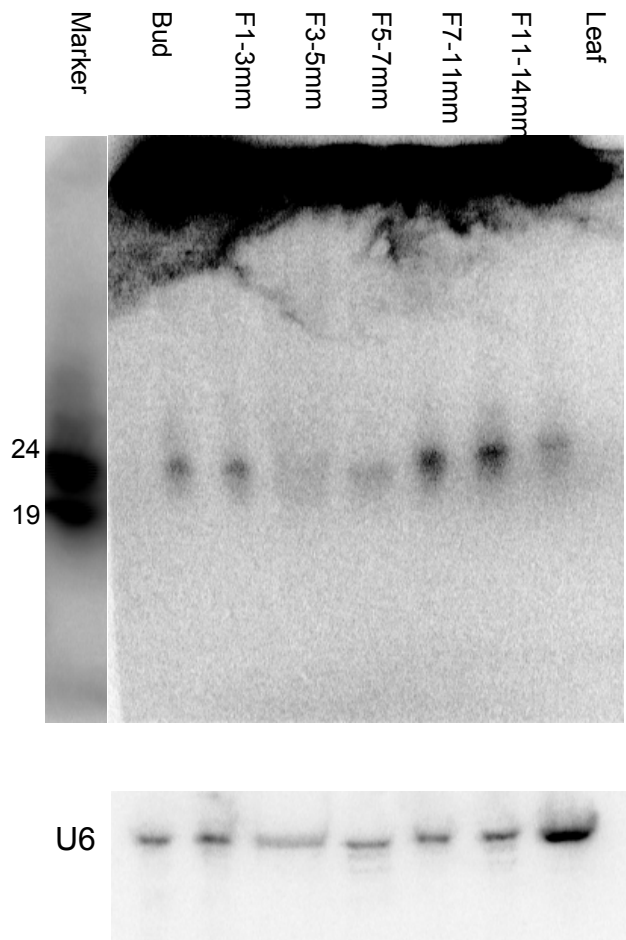
Tom193



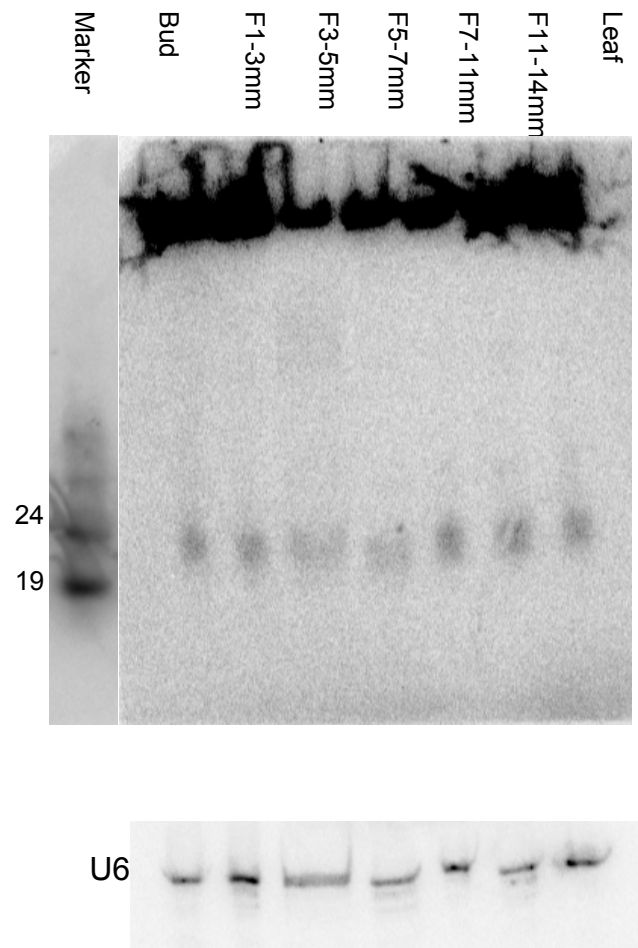
Tom32



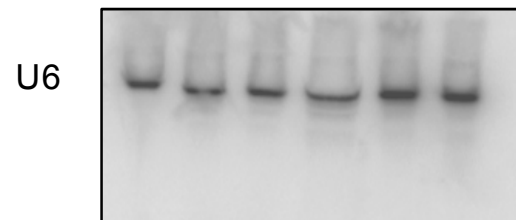
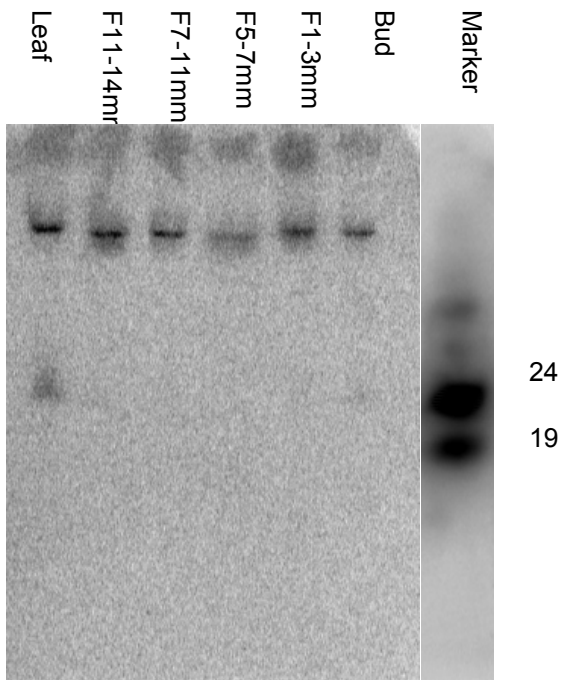
Tom40



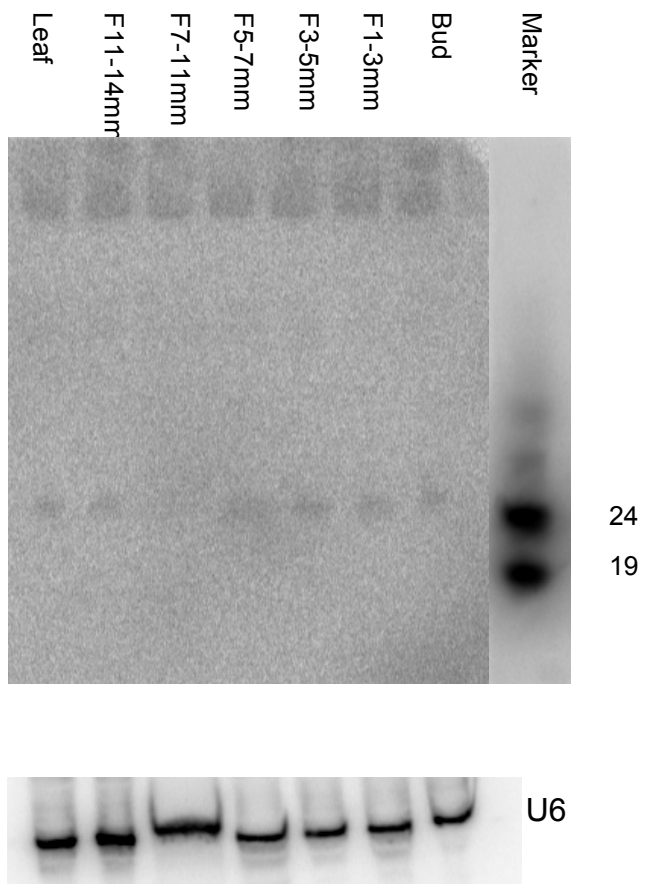
Tom44



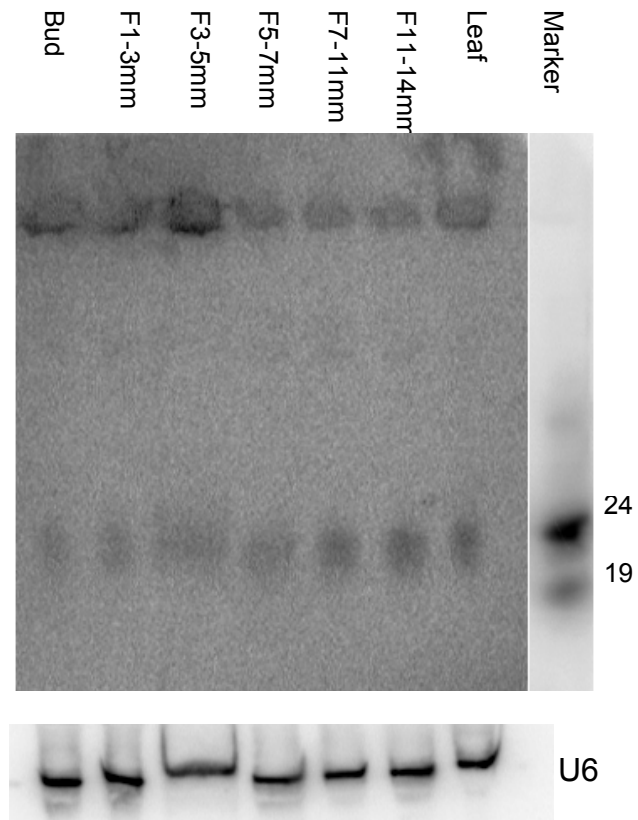
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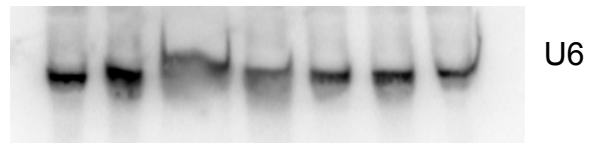
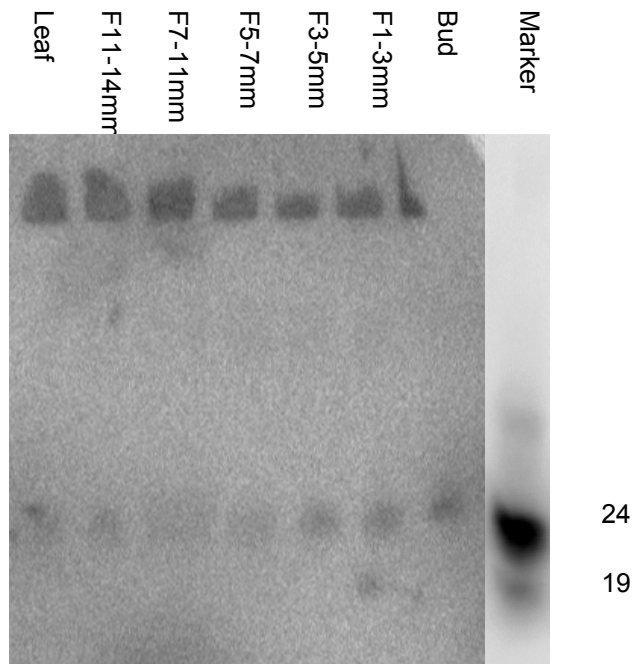
Tom82



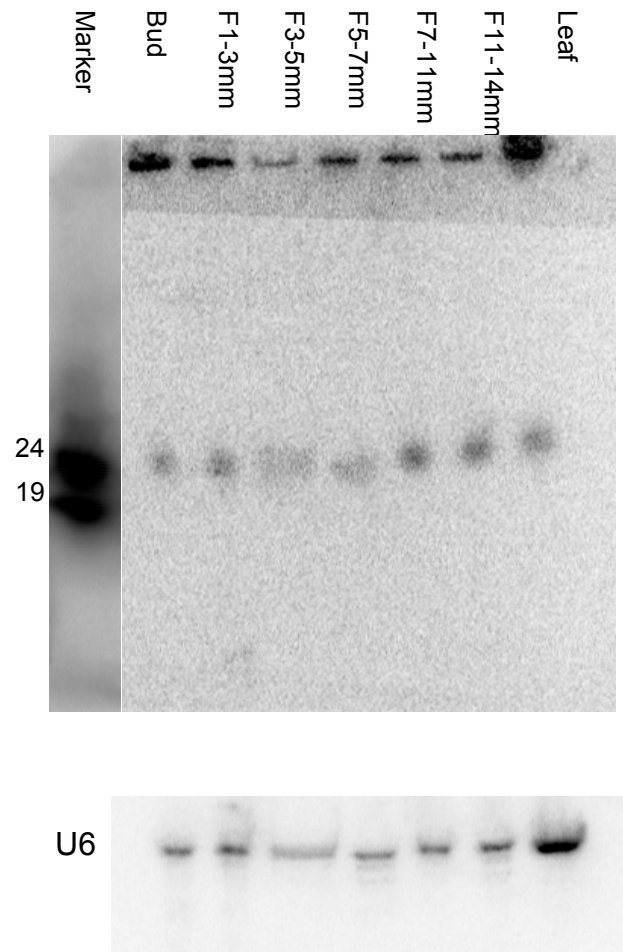
Tom153



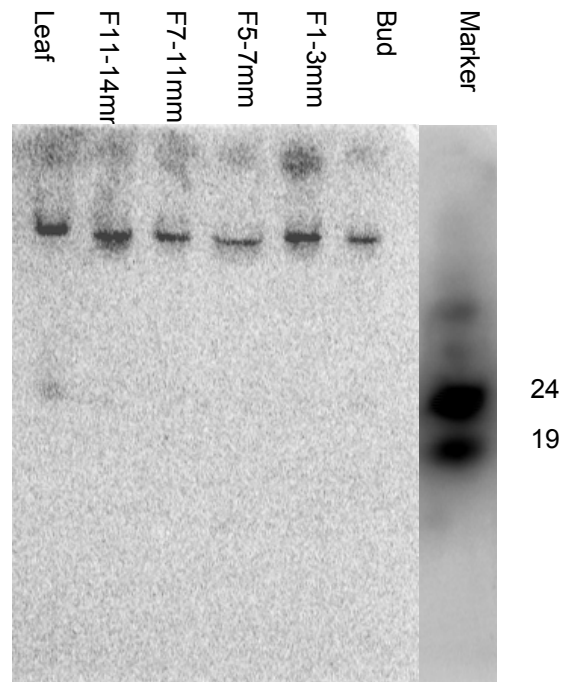
Tom101



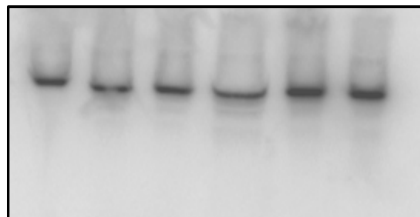
Tom108



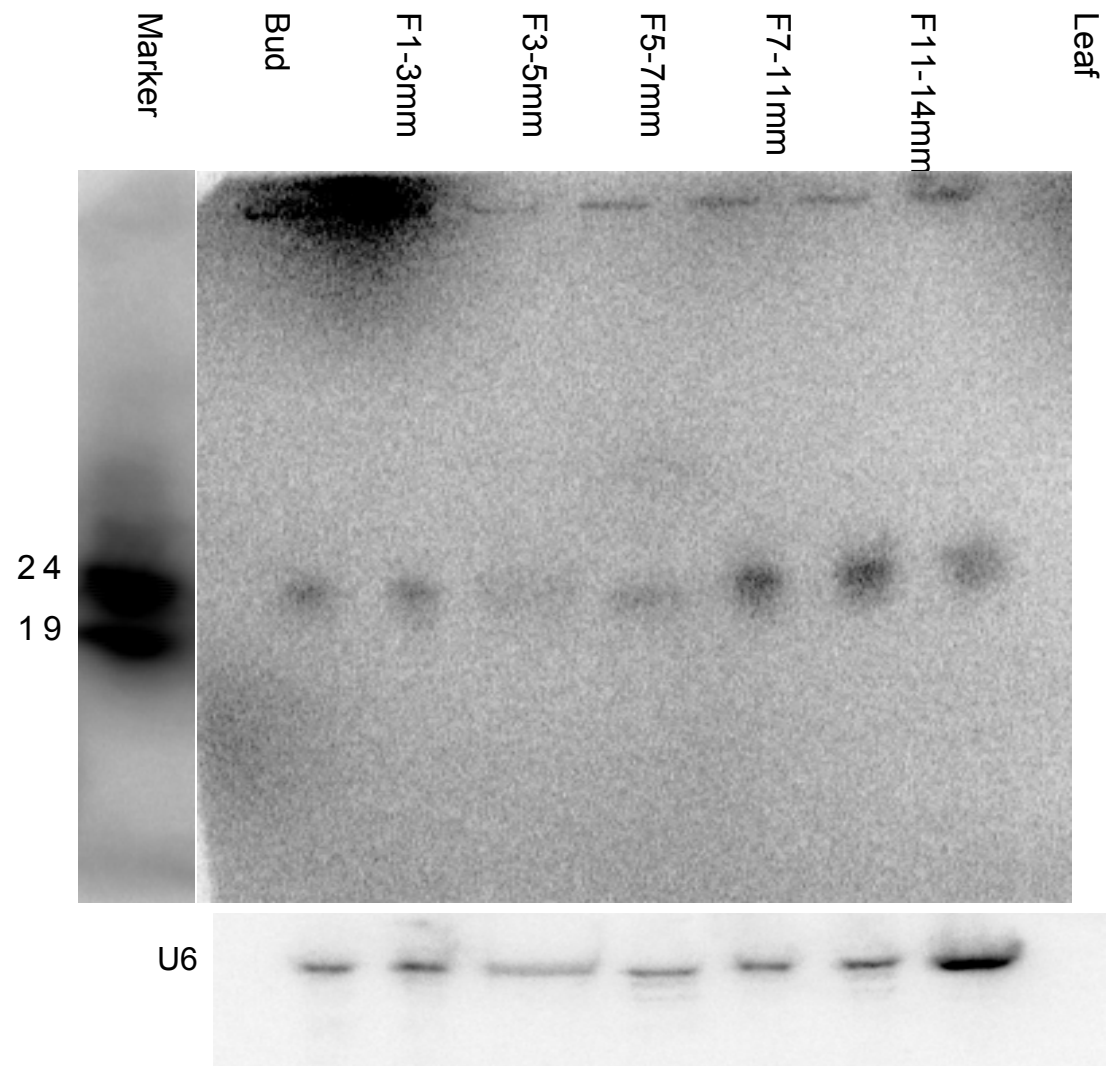
Tom154



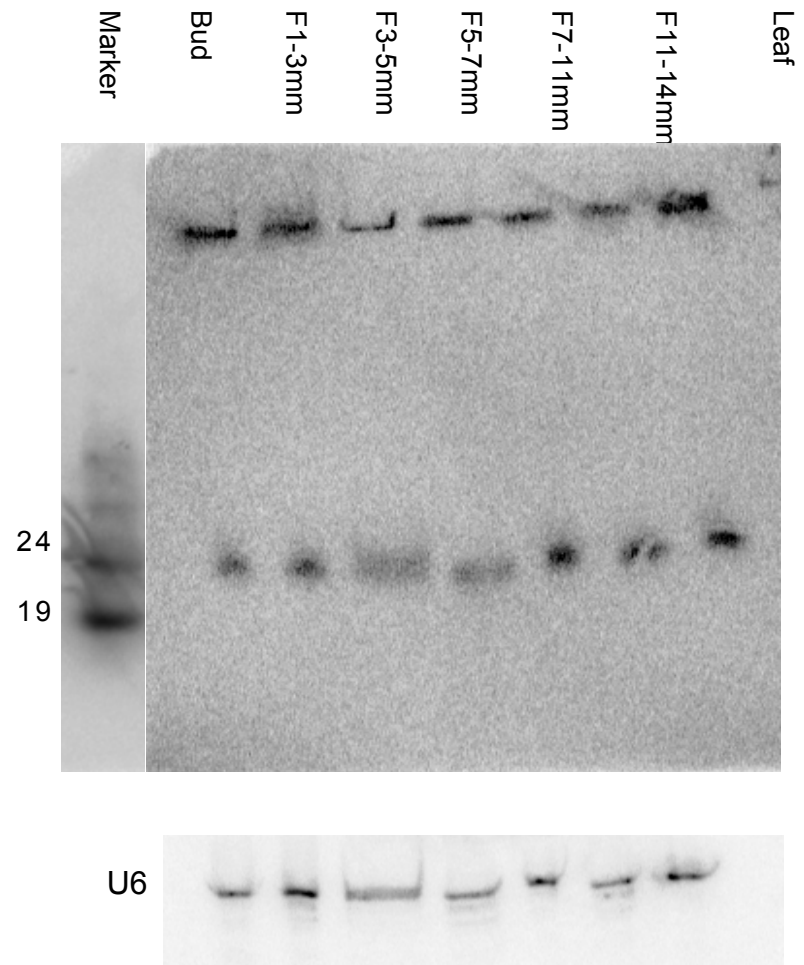
U6



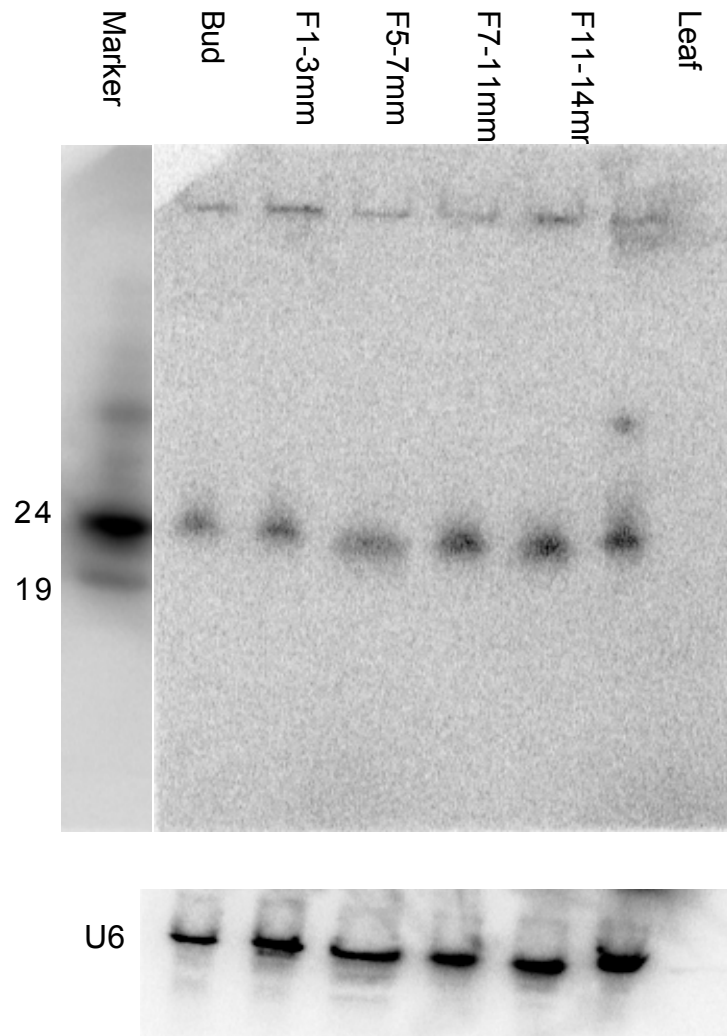
Tom156



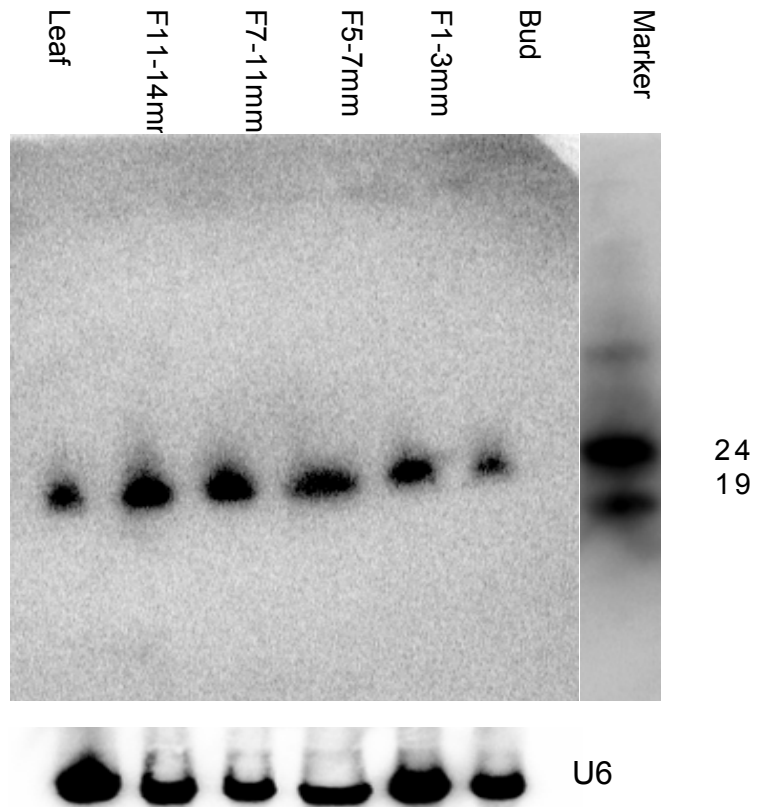
Tom160



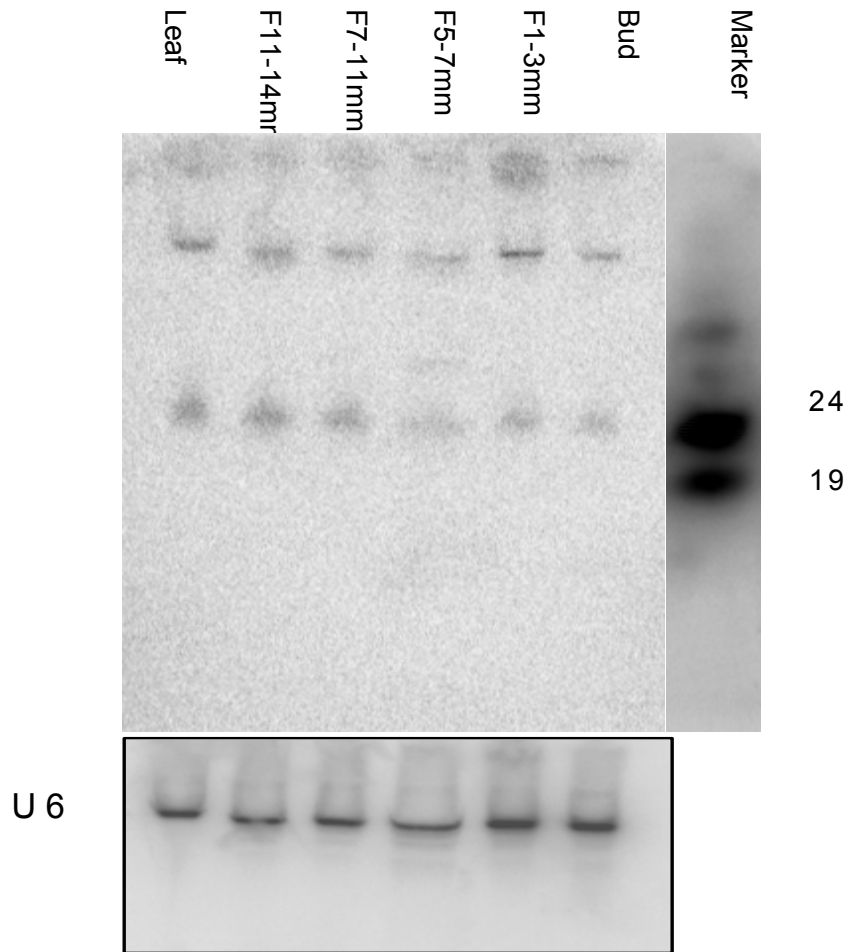
Tom175



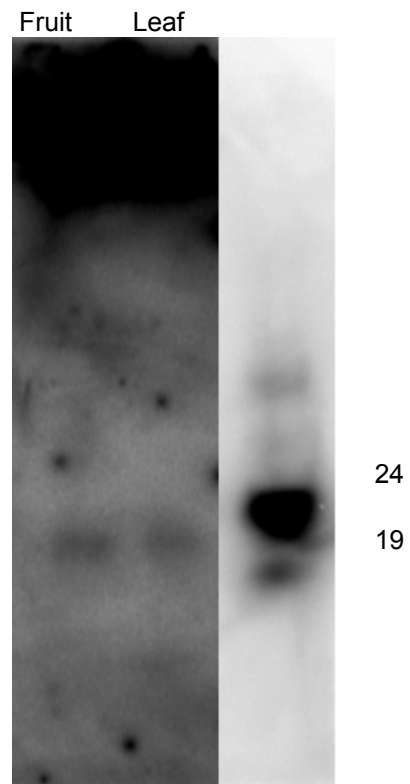
Tom189



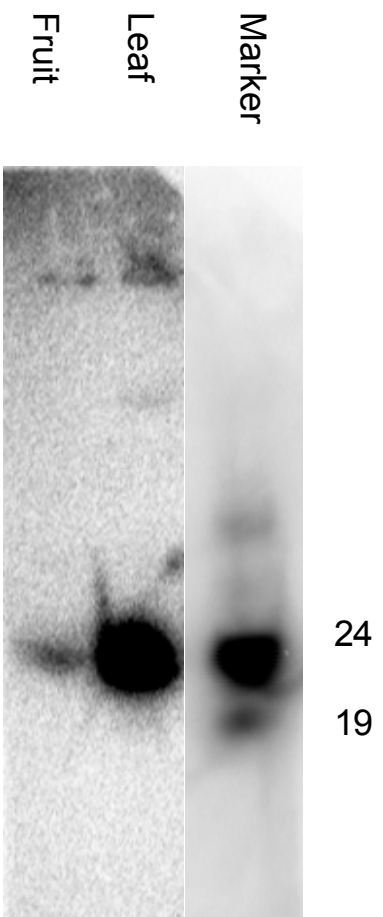
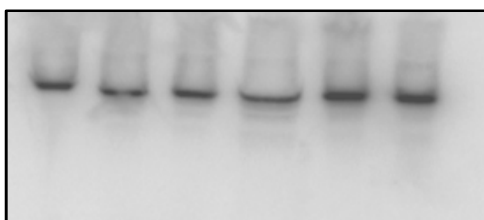
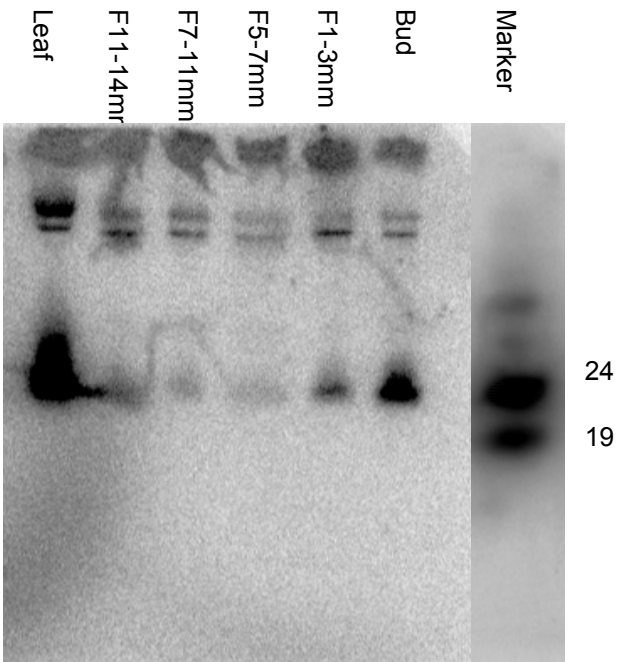
Tom191



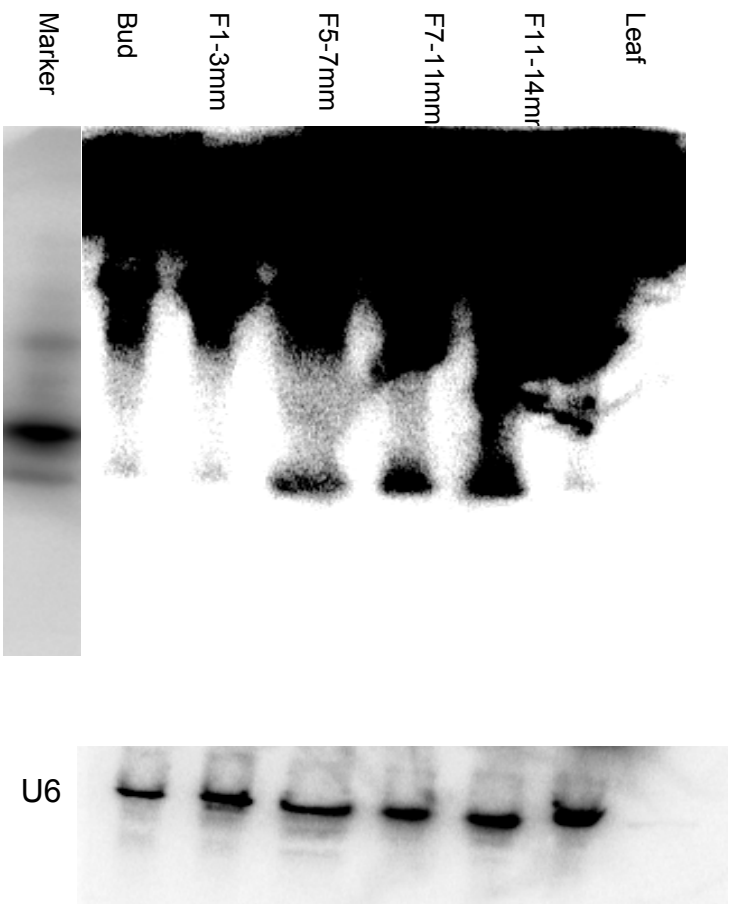
NGM1



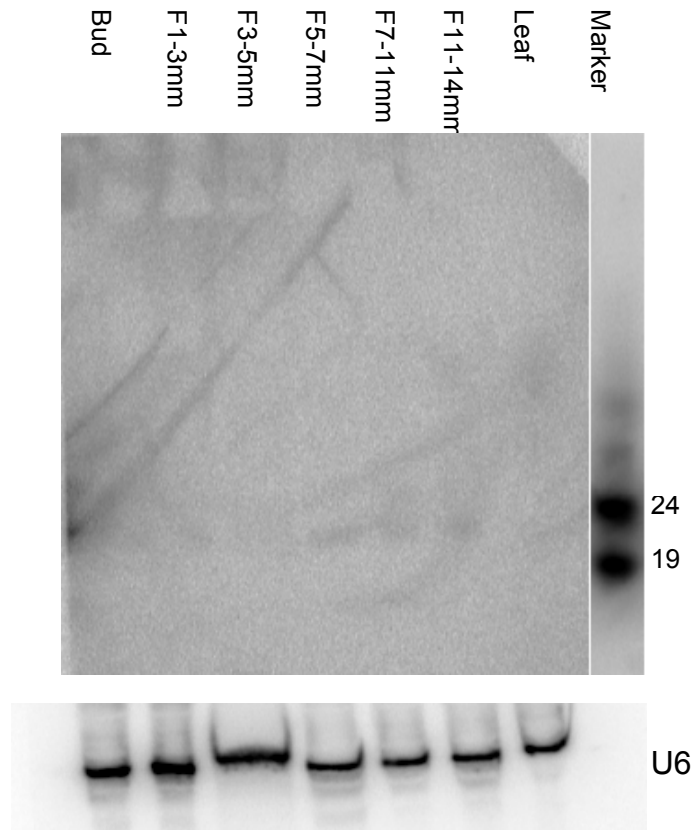
NGM3



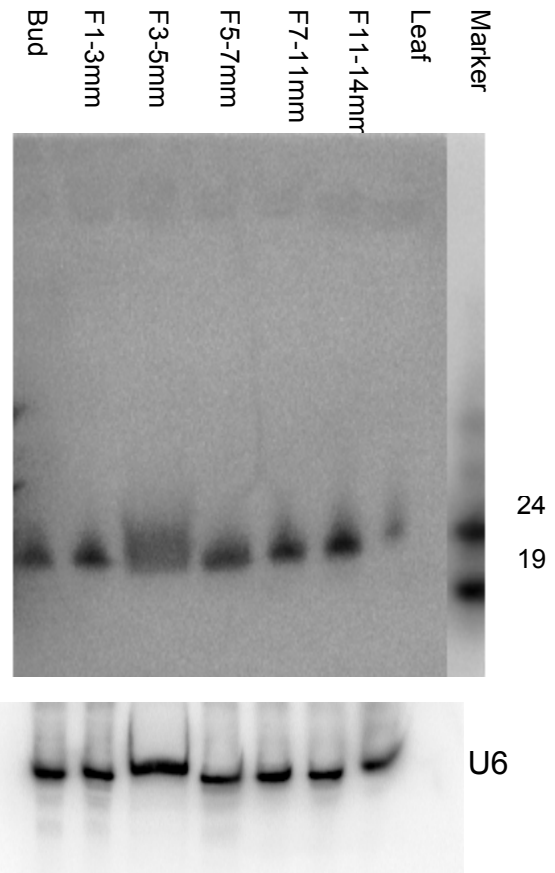
NGM6



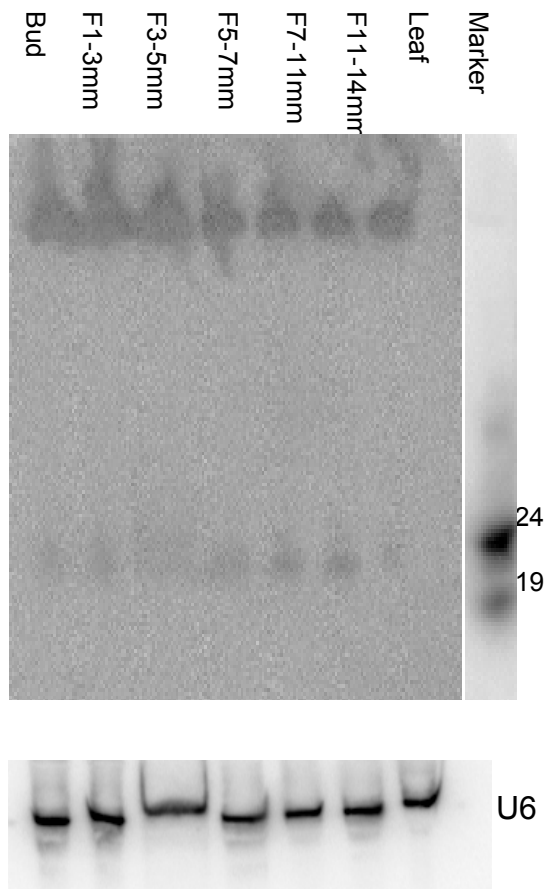
Top1



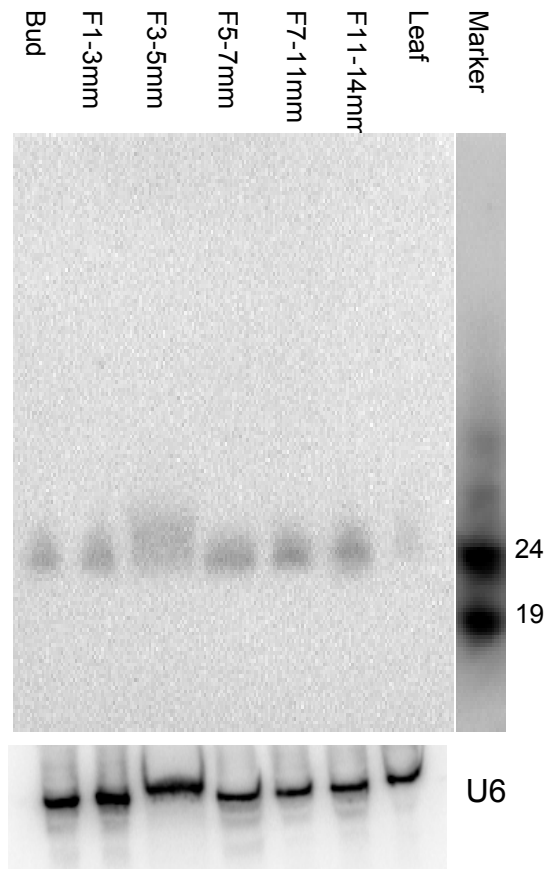
Top2



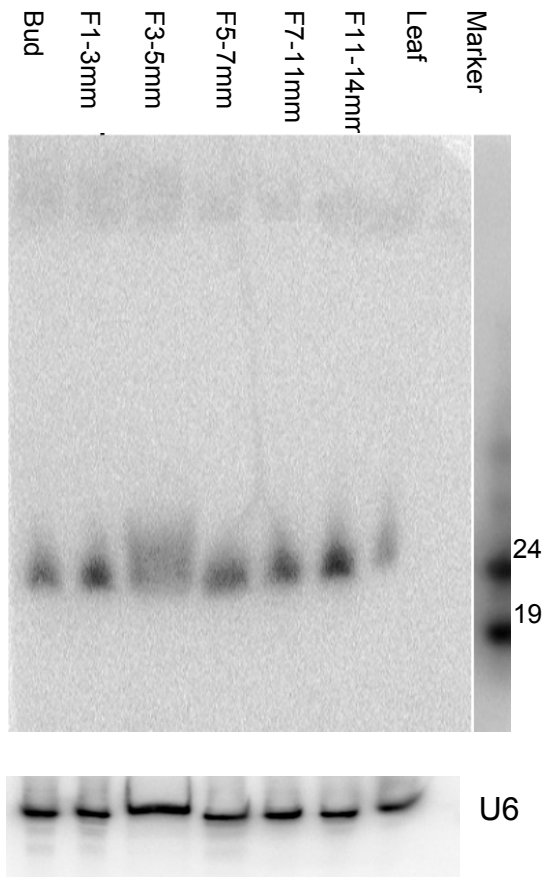
Top3



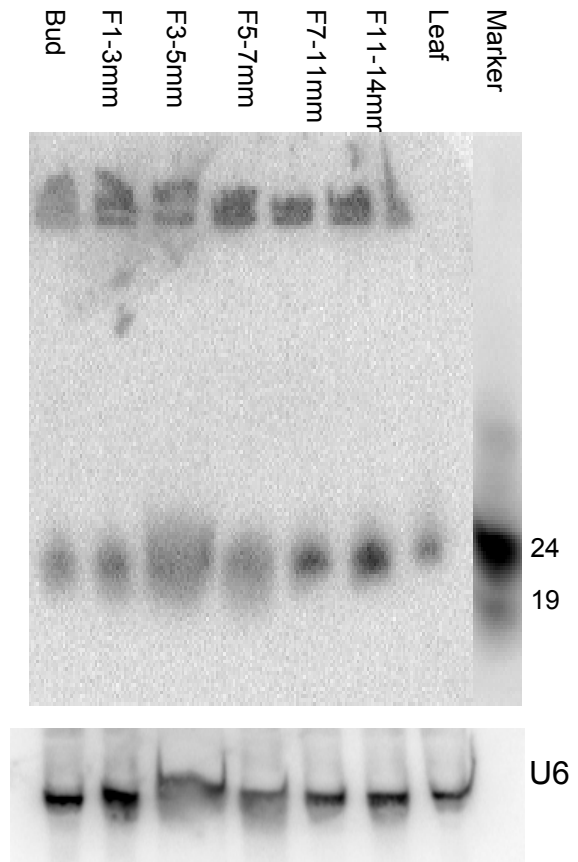
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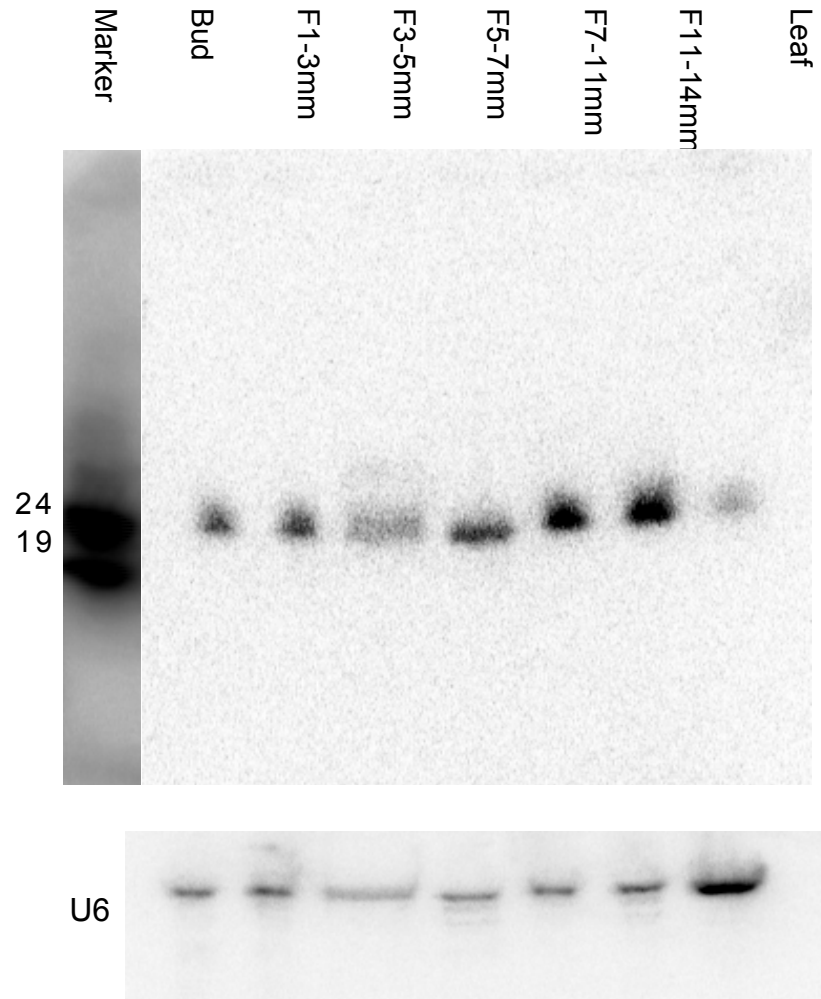
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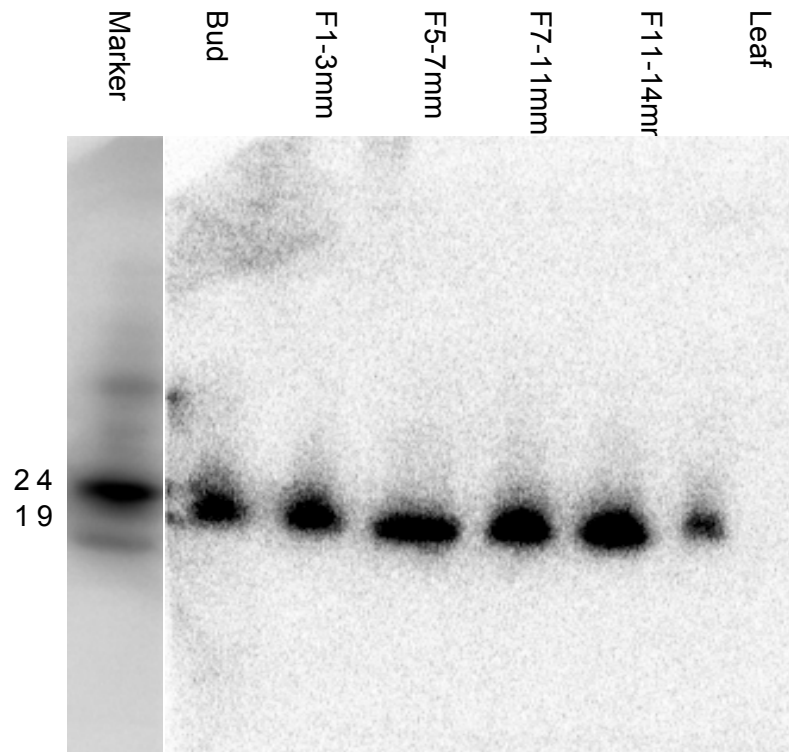
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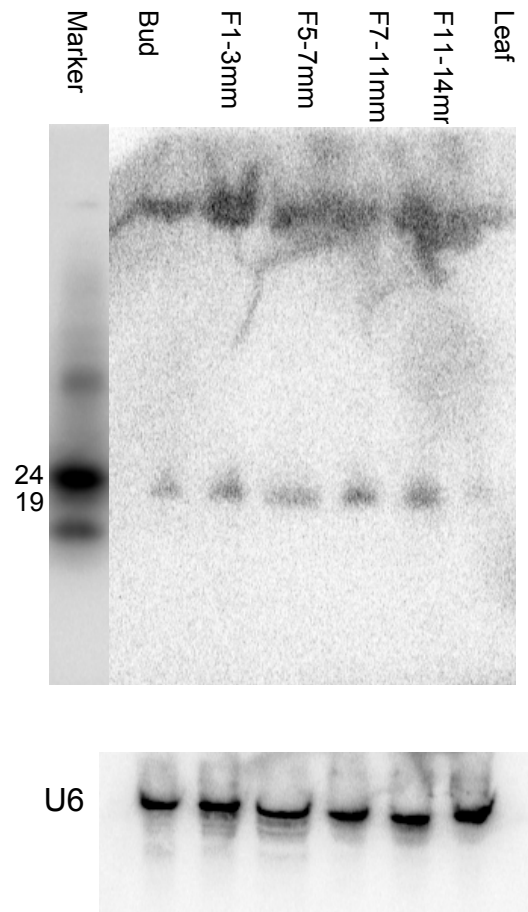
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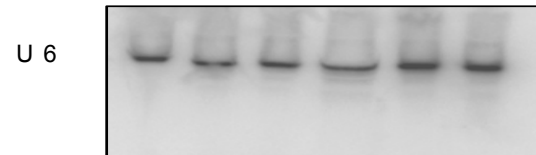
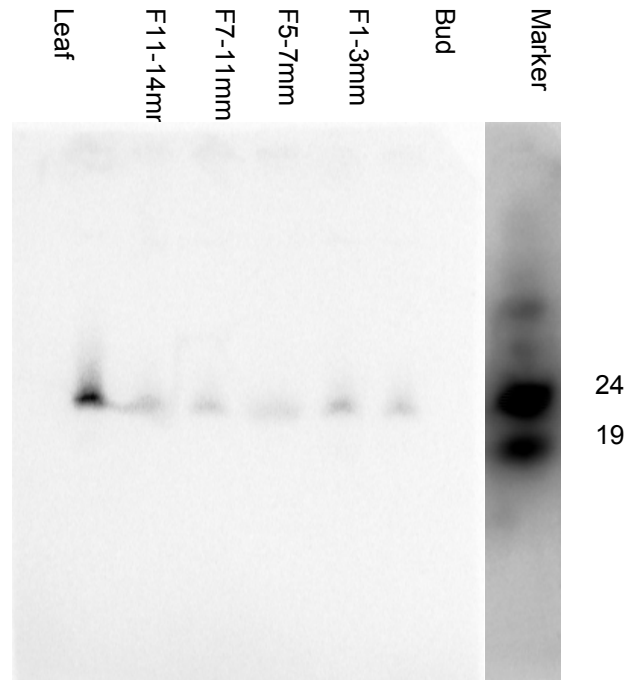
Top11



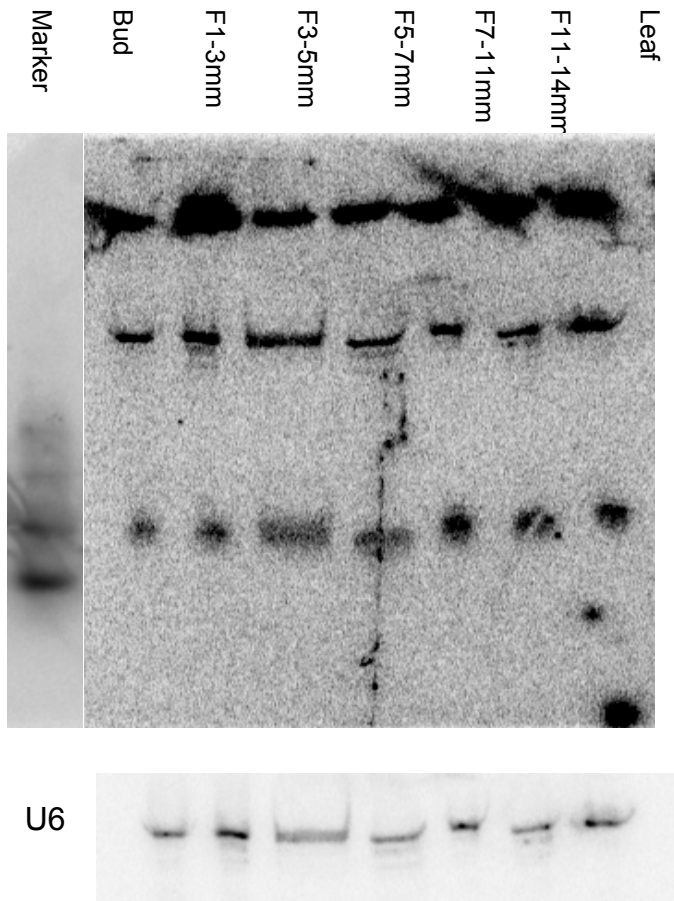
Top12



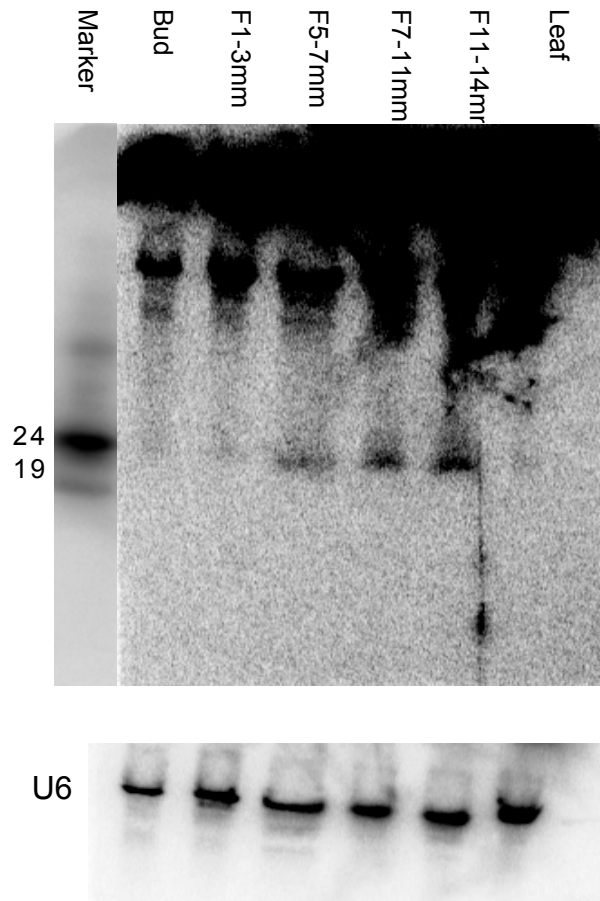
Top15



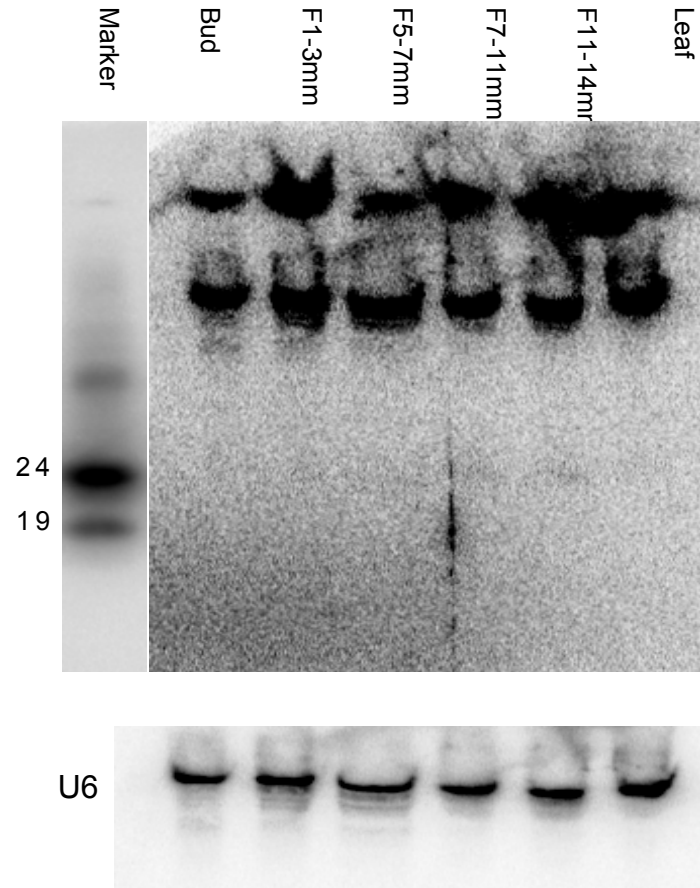
Tom Tar2



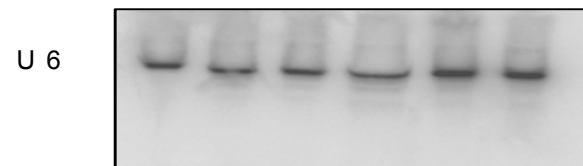
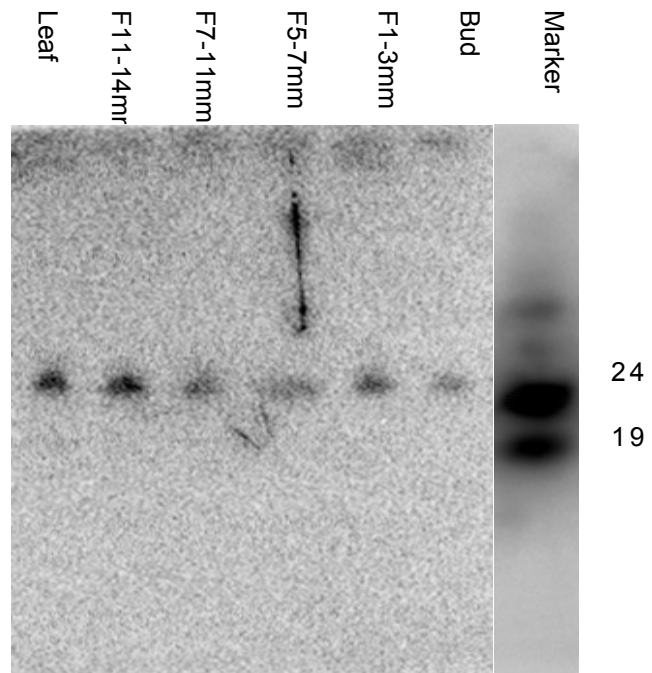
Tom Tar3



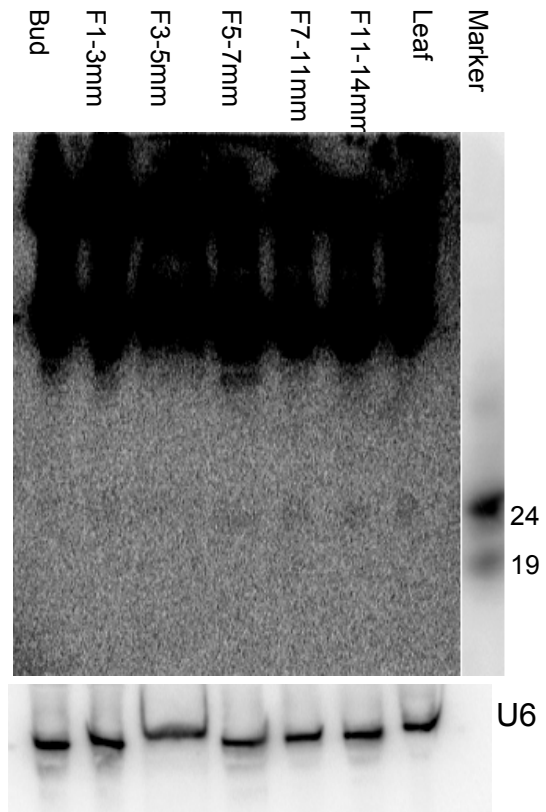
Tom Tar4



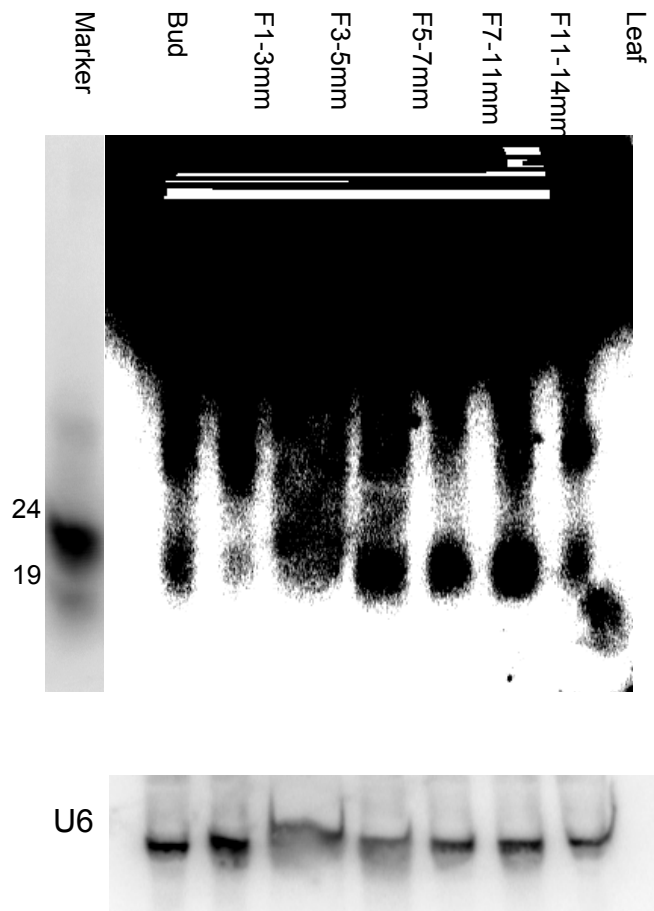
Tom Tar6



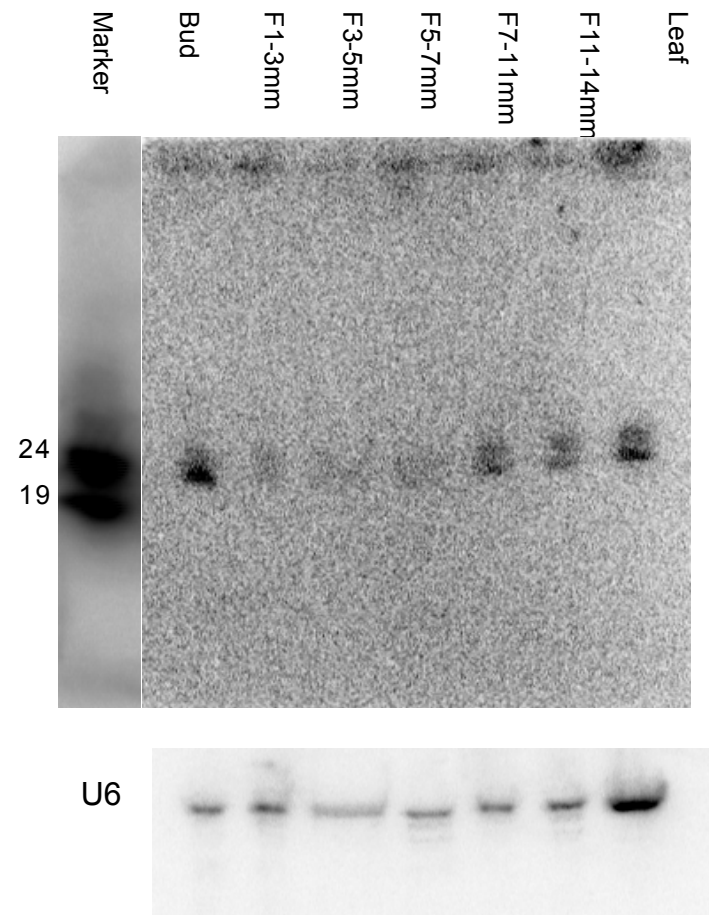
Tom Tar9



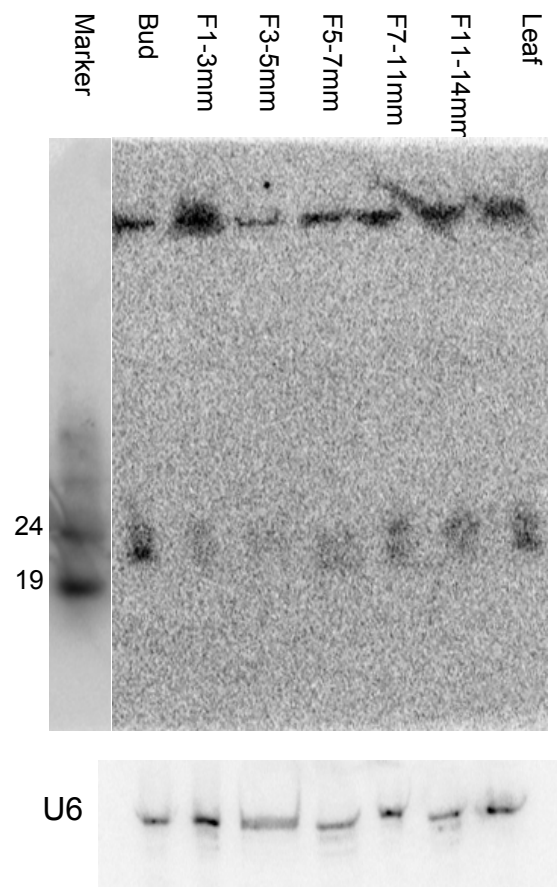
Tom Tar10



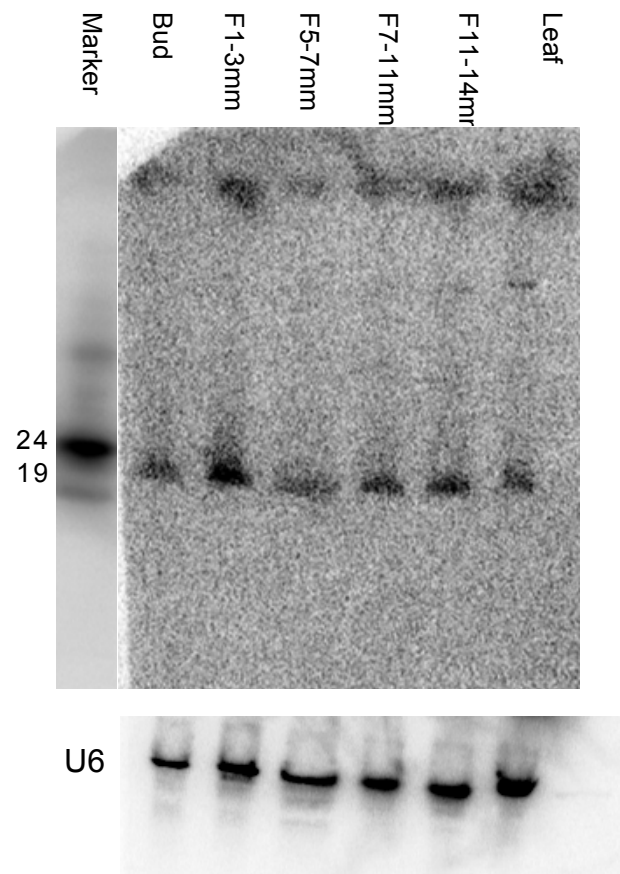
Tom Tar11



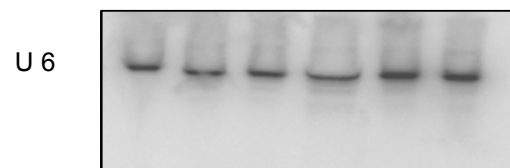
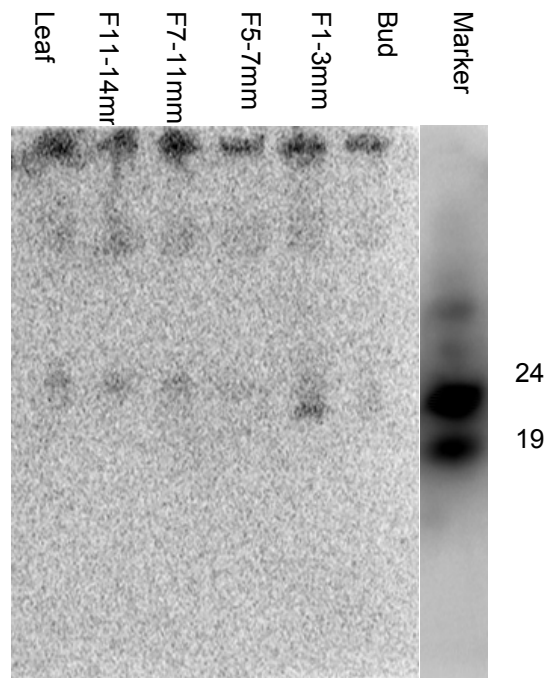
Tom Tar12



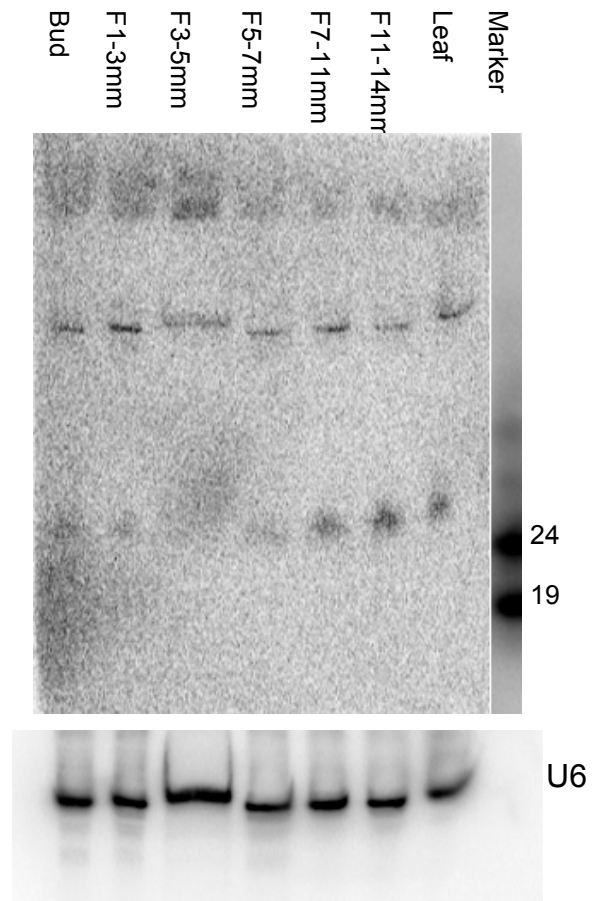
Tom Tar13



Tom Tar16



Tom Tar18



Tom Tar19

